

## Claims

1. A method for the detection of drink-spoiling microorganisms in a sample, whereby the detection is carried out by using at least one oligonucleotide probe  
5 having a nucleic acid sequence selected from the group consisting of (all sequences in 5' → 3' direction):

	SEQ ID No. 1:	5'- GTTTGACCAGATTCTCCGCTC
	SEQ ID No. 5:	5'- CCCGGTCGAATTAAAACC
10	SEQ ID No. 6:	5'- GCCCGGTCGAATTAAAAC
	SEQ ID No. 7:	5'- GGCCCGGTCGAATTAAAA
	SEQ ID No. 8:	5'- AGGCCCGGTCGAATTAAA
	SEQ ID No. 9:	5'- AAGGCCCGGTCGAATTAA
	SEQ ID No. 10:	5'- ATATTTCGAGCGAAACGCC
15	SEQ ID No. 11:	5'- AAAGATCCGGACCGGCCG
	SEQ ID No. 12	5'- GGAAAGATCCGGACCGGC
	SEQ ID No. 13	5'- GAAAGATCCGGACCGGCC
	SEQ ID No. 14	5'- GATCCGGACCGGCCGACC
	SEQ ID No. 15	5'- AGATCCGGACCGGCCGAC
20	SEQ ID No. 16	5'- AAGATCCGGACCGGCCGA
	SEQ ID No. 17	5'- GAAAGGCCCGGTCGAATT
	SEQ ID No. 18	5'- AAAGGCCCGGTCGAATTA
	SEQ ID No. 19	5'- GGAAAGGCCCGGTCGAAT
	SEQ ID No. 20	5'- AGGAAAGGCCCGGTCGAA
25	SEQ ID No. 21	5'- AAGGAAAGGCCCGGTCGA
	SEQ ID No. 22:	5'- ATAGCACTGGGATCCTCGCC
	SEQ ID No. 23:	5'- CCAGCCCCAAAGTTACCTTC
	SEQ ID No. 24:	5'- TCCTTGACGTAAAGTCGCAG
	SEQ ID No. 25:	5'- GGAAGAAAACCAGTACGC
30	SEQ ID No. 26:	5'- CCGGTCGGAAGAAAACCA

	SEQ ID No. 27:	5'- GAAGAAAACCAAGTACGCG
	SEQ ID No. 28:	5'- CCCGGTCGGAAGAAAACC
	SEQ ID No. 29:	5'- CGGTCGGAAGAAAACCAAG
	SEQ ID No. 30:	5'- GGTCGGAAGAAAACCAAGT
5	SEQ ID No. 31:	5'- AAGAAAACCAAGTACGCGG
	SEQ ID No. 32:	5'- GTACGCGGA AAAAATCCGG
	SEQ ID No. 33:	5'- AGTACGCGGA AAAAATCCG
	SEQ ID No. 34:	5'- GCGGA AAAAATCCGGACCG
	SEQ ID No. 35:	5'- CGGAAGAAAACCAAGTACG
10	SEQ ID No. 36:	5'- GCCCGGTCGGAAGAAAAC
	SEQ ID No. 37:	5'- CGCGGA AAAAATCCGGACC
	SEQ ID No. 38:	5'- CAGTACGCGGA AAAAATCC
	SEQ ID No. 39:	5'- AGAAAACCAAGTACGCGGA
	SEQ ID No. 40:	5'- GGCCCGGTCGGAAGAAAA
15	SEQ ID No. 41:	5'- ATAAACACCACCCGATCC
	SEQ ID No. 42:	5'- ACGCGGA AAAAATCCGGAC
	SEQ ID No. 43:	5'- GAGAGGCCCGGTCGGAAG
	SEQ ID No. 44:	5'- AGAGGCCCGGTCGGAAGA
	SEQ ID No. 45:	5'- GAGGCCCGGTCGGAAGAA
20	SEQ ID No. 46:	5'- AGGCCCGGTCGGAAGAAA
	SEQ ID No. 47:	5'- CCGAGTGGGTCAGTAAAT
	SEQ ID No. 48:	5'- CCAGTACGCGGA AAAAATC
	SEQ ID No. 49:	5'- TAAACACCACCCGATCCC
	SEQ ID No. 50:	5'- GGAGAGGCCCGGTCGGAA
25	SEQ ID No. 51:	5'- GAAAACCAAGTACGCGGAA
	SEQ ID No. 52:	5'- TACGCGGA AAAAATCCGGA
	SEQ ID No. 53:	5'- GGCCACAGGGACCCAGGG
	SEQ ID No. 54:	5'- TCACCAAGGGCCACAGGG
	SEQ ID No. 55:	5'- GGGCCACAGGGACCCAGG
30	SEQ ID No. 56:	5'- TTCACCAAGGGCCACAGG

	SEQ ID No. 57:	5'- ACAGGGACCCAGGGCTAG
	SEQ ID No. 58:	5'- AGGGCCACAGGGACCCAG
	SEQ ID No. 59:	5'- GTTCACCAAGGGCCACAG
	SEQ ID No. 60:	5'- GCCACAGGGACCCAGGGC
5	SEQ ID No. 61:	5'- CAGGGACCCAGGGCTAGC
	SEQ ID No. 62:	5'- AGGGACCCAGGGCTAGCC
	SEQ ID No. 63:	5'- ACCAAGGGCCACAGGGAC
	SEQ ID No. 64:	5'- CCACAGGGACCCAGGGCT
	SEQ ID No. 65:	5'- CACAGGGACCCAGGGCTA
10	SEQ ID No. 66:	5'- CACCAAGGGCCACAGGGA
	SEQ ID No. 67:	5'- GGGACCCAGGGCTAGCCA
	SEQ ID No. 68:	5'- AGGAGAGGCCCGGTCGGA
	SEQ ID No. 69:	5'- AAGGAGAGGCCCGGTCGG
	SEQ ID No. 70:	5'- GAAGGAGAGGCCCGGTCG
15	SEQ ID No. 71:	5'- AGGGCTAGCCAGAAGGAG
	SEQ ID No. 72:	5'- GGGCTAGCCAGAAGGAGA
	SEQ ID No. 73:	5'- AGAAGGAGAGGCCCGGTC
	SEQ ID No. 74:	5'- CAAGGGCCACAGGGACCC
	SEQ ID No. 75:	5'- CCAAGGGCCACAGGGACC
20	SEQ ID No. 76:	5'- GTCGGAAAAACCAAGTACG
	SEQ ID No. 77:	5'- GCCCGGTCGGAAAAACCA
	SEQ ID No. 78:	5'- CCGGTCGGAAAAACCAAGT
	SEQ ID No. 79:	5'- CCCGGTCGGAAAAACCAAG
	SEQ ID No. 80:	5'- TCGGAAAAACCAAGTACGC
25	SEQ ID No. 81:	5'- CGGAAAAACCAAGTACGCG
	SEQ ID No. 82:	5'- GGAAAAACCAAGTACGCGG
	SEQ ID No. 83:	5'- GTACGCGGAAAAATCCGG
	SEQ ID No. 84:	5'- AGTACGCGGAAAAATCCG
	SEQ ID No. 85:	5'- GCGGAAAAATCCGGACCG
30	SEQ ID No. 86:	5'- GGTCGGAAAAACCAAGTAC

	SEQ ID No. 87:	5'- ACTCCTAGTGGTGCCCTT
	SEQ ID No. 88:	5'- GCTCCACTCCTAGTGGTG
	SEQ ID No. 89:	5'- CACTCCTAGTGGTGCCCT
	SEQ ID No. 90:	5'- CTCCACTCCTAGTGGTGC
5	SEQ ID No. 91:	5'- TCCACTCCTAGTGGTGCC
	SEQ ID No. 92:	5'- CCACTCCTAGTGGTGCCC
	SEQ ID No. 93:	5'- GGCTCCACTCCTAGTGGT
	SEQ ID No. 94:	5'- AGGCTCCACTCCTAGTGG
	SEQ ID No. 95:	5'- GGCCCGGTCGGAAAAACC
10	SEQ ID No. 96:	5'- GAAAAACCAAGTACGCGGA
	SEQ ID No. 97:	5'- CGCGGAAAAATCCGGACC
	SEQ ID No. 98:	5'- CAGTACGCGGAAAAATCC
	SEQ ID No. 99:	5'- CGGTCGGAAAAACCAAGTA
	SEQ ID No. 100:	5'- AAGGCCCGGTCGGAAAAA
15	SEQ ID No. 101:	5'- CAGGCTCCACTCCTAGTG
	SEQ ID No. 102:	5'- CTCCTAGTGGTGCCCTTC
	SEQ ID No. 103:	5'- TCCTAGTGGTGCCCTTCC
	SEQ ID No. 104:	5'- GCAGGCTCCACTCCTAGT
	SEQ ID No. 105:	5'- AGGCCCGGTCGGAAAAAC
20	SEQ ID No. 106:	5'- ACGCGGAAAAATCCGGAC
	SEQ ID No. 107:	5'- CCAGTACGCGGAAAAATC
	SEQ ID No. 108:	5'- CTAGTGGTGCCCTTCGCT
	SEQ ID No. 109:	5'- GAAAGGCCCGGTCGGAAA
	SEQ ID No. 110:	5'- AAAGGCCCGGTCGGAAAA
25	SEQ ID No. 111:	5'- TACGCGGAAAAATCCGGA
	SEQ ID No. 112:	5'- GGAAAGGCCCGGTCGGAA
	SEQ ID No. 113:	5'- ATCTCTTCCGAAAGGTCG
	SEQ ID No. 114:	5'- CATCTCTTCCGAAAGGTC
	SEQ ID No. 115:	5'- CTCTTCCGAAAGGTCGAG
30	SEQ ID No. 116:	5'- CTTCCGAAAGGTCGAGAT

	SEQ ID No. 117:	5'- TCTCTTCCGAAAGGTCGA
	SEQ ID No. 118:	5'- TCTTCCGAAAGGTCGAGA
	SEQ ID No. 119:	5'- CCTAGTGGTGCCCTTCCG
	SEQ ID No. 120:	5'- TAGTGGTGCCCTTCCGTC
5	SEQ ID No. 121:	5'- AGTGGTGCCCTTCCGTCA
	SEQ ID No. 122:	5'- GCCAAGGTTAGACTCGTT
	SEQ ID No. 123:	5'- GGCCAAGGTTAGACTCGT
	SEQ ID No. 124:	5'- CCAAGGTTAGACTCGTTG
	SEQ ID No. 125:	5'- CAAGGTTAGACTCGTTGG
10	SEQ ID No. 126:	5'- AAGGTTAGACTCGTTGGC
	SEQ ID No. 127:	5'- CTCGCCTCACGGGGTTCTCA
	SEQ ID No. 128:	5'- GGCCCGGTCGAAATTAAA
	SEQ ID No. 129:	5'- AGGCCCGGTCGAAATTAA
	SEQ ID No. 130:	5'- AAGGCCCGGTCGAAATTA
15	SEQ ID No. 131:	5'- AAAGGCCCGGTCGAAATT
	SEQ ID No. 132:	5'- GAAAGGCCCGGTCGAAAT
	SEQ ID No. 133:	5'- ATATTCGAGCGAAACGCC
	SEQ ID No. 134:	5'- GGAAAGGCCCGGTCGAAA
	SEQ ID No. 135:	5'- AAAGATCCGGACCGGCCG
20	SEQ ID No. 136:	5'- GGAAAGATCCGGACCGGC
	SEQ ID No. 137:	5'- GAAAGATCCGGACCGGCC
	SEQ ID No. 138:	5'- GATCCGGACCGGECGAEC
	SEQ ID No. 139:	5'- AGATCCGGACCGGCCGAC
	SEQ ID No. 140:	5'- AAGATCCGGACCGGCCGA
25	SEQ ID No. 141:	5'- AGGAAAGGCCCGGTCGAA
	SEQ ID No. 142:	5'- AAGGAAAGGCCCGGTCGA
	SEQ ID No. 143:	5'-CGAGCAAAACGCCTGCTTTG
	SEQ ID No. 144:	5'-CGCTCTGAAAGAGAGTTGCC
	SEQ ID No. 145:	5'-AGTTGCCCCCTACACTAGAC..
30	SEQ ID No. 146:	5'-GCTTCTCCGTCCCGCGCCG

SEQ ID No. 148: 5'- CCTGGTTCGCCAAAAAGGC  
SEQ ID No. 149: 5'-GATTCTCGGCCCCATGGG  
SEQ ID No. 150: 5'- ACCCTCTACGGCAGCCTGTT  
SEQ ID No. 151: 5'- GATCGGTCTCCAGCGATTCA  
5 SEQ ID No. 152: 5'- ACCCTCCACGGCGGCCTGTT  
SEQ ID No. 153: 5'- GATTCTCCGCGCCATGGG  
SEQ ID No. 154: 5'- TCATCAGACGGGATTCTCAC  
SEQ ID No. 157: 5'-AGTTGCCCCCTCCTCTAAGC  
SEQ ID No. 158: 5'-CTGCCACAAGGACAAATGGT  
10 SEQ ID No. 159: 5'-TGCCCCCTCTTCTAAGCAAAT  
SEQ ID No. 160: 5'-CCCCAAAGTTGCCCTCTC  
SEQ ID No. 163: 5'-AAGACCAGGCCACCTCAT  
SEQ ID No. 164: 5'- CATCATAGAACACCGTCC  
SEQ ID No. 165: 5'- CCTTCCGAAGTCGAGGTTTT  
15 SEQ ID No. 166: 5'- GGGAGTGTTGCCAACTC  
SEQ ID No. 167: 5'- AGCGGTCGTTCGCAACCCT  
SEQ ID No. 168: 5'- CCGAAGTCGGGGTTTTGCGG  
SEQ ID No. 169: 5'- GATAGCCGAAACCACCTTTC  
SEQ ID No. 170: 5'- GCCGAAACCACCTTTCAAAC  
20 SEQ ID No. 171: 5'- GTGATAGCCGAAACCACCTT  
SEQ ID No. 172: 5'- AGTGATAGCCGAAACCACCT  
SEQ ID No. 173: 5'- TTTAACGGGATGCGTTCGAG  
SEQ ID No. 174: 5'- AAGTGATAGCCGAAACCACC  
SEQ ID No. 175: 5'- GGTTGAATACCGTCAACGTC  
25 SEQ ID No. 176: 5'- GCACAGTATGTCAAGACCTG  
SEQ ID No. 177: 5'- CATCCGATGTGCAAGCACTT  
SEQ ID No. 178: 5'- TCATCCGATGTGCAAGCACT  
SEQ ID No. 179: 5'- CCGATGTGCAAGCACTTCAT  
SEQ ID No. 180: 5'- CCACTCATCCGATGTGCAAG  
30 SEQ ID No. 181: 5'- GCCACAGTTCGCCACTCATC

	SEQ ID No. 182:	5'- CCTCCGCGTTTGTCAACGGC
	SEQ ID No. 183:	5'- ACCAGTTCGCCACAGTTCGC
	SEQ ID No. 184:	5'- CACTCATCCGATGTGCAAGC
	SEQ ID No. 185:	5'- CCAGTTCGCCACAGTTCGCC
5	SEQ ID No. 186:	5'- CTCATCCGATGTGCAAGCAC
	SEQ ID No. 187:	5'- TCCGATGTGCAAGCACTTCA
	SEQ ID No. 188:	5'- CGCCACTCATCCGATGTGCA
	SEQ ID No. 189:	5'- CAGTTCGCCACAGTTCGCCA
	SEQ ID No. 190:	5'- GCCACTCATCCGATGTGCAA
10	SEQ ID No. 191:	5'- CGCCACAGTTCGCCACTCAT
	SEQ ID No. 192:	5'- ATCCGATGTGCAAGCACTTC
	SEQ ID No. 193:	5'- GTTCGCCACAGTTCGCCACT
	SEQ ID No. 194:	5'- TCCTCCGCGTTTGTCAACGG
	SEQ ID No. 195:	5'- CGCCAGGGTTCATCCTGAGC
15	SEQ ID No. 196:	5'- AGTTCGCCACAGTTCGCCAC
	SEQ ID No. 197:	5'- TCGCCACAGTTCGCCACTCA
	SEQ ID No. 198:	5'- TTAACGGGATGCGTTCGACT
	SEQ ID No. 199:	5'- TCGCCACTCATCCGATGTGC
	SEQ ID No. 200:	5'- CCACAGTTCGCCACTCATCC
20	SEQ ID No. 201:	5'- GATTTAACGGGATGCGTTTCG
	SEQ ID No. 202:	5'- TAACGGGATGCGTTCGACTT ---
	SEQ ID No. 203:	5'- AACGGGATGCGTTCGACTTG ---
	SEQ ID No. 204:	5'- CGAAGGTTACCGAACCGACT
	SEQ ID No. 205:	5'- CCGAAGGTTACCGAACCGAC
25	SEQ ID No. 206:	5'- CCCGAAGGTTACCGAACCGA
	SEQ ID No. 207:	5'- TTCCTCCGCGTTTGTCAACCG
	SEQ ID No. 208:	5'- CCGCCAGGGTTCATCCTGAG
	SEQ ID No. 209:	5'- TCCTTCCAGAAGTGATAGCC
	SEQ ID No. 210:	5'- CACCAGTTCGCCACAGTTCG
30	SEQ ID No. 211:	5'- ACGGGATGCGTTCGACTTGC ---

	SEQ ID No. 212:	5'- GTCCTTCCAGAAGTGATAGC
	SEQ ID No. 213:	5'- GCCAGGGTTCATCCTGAGCC
	SEQ ID No. 214:	5'- ACTCATCCGATGTGCAAGCA
	SEQ ID No. 215:	5'- ATCATTGCCTTGGTGAACCG
5	SEQ ID No. 216:	5'- TCCGCGTTTGTACACCGGCAG
	SEQ ID No. 217:	5'- TGAACCGTTACTCCACCAAC
	SEQ ID No. 218:	5'- GAAGTGATAGCCGAAACCAC
	SEQ ID No. 219:	5'- CCGCGTTTGTACACCGGCAGT
	SEQ ID No. 220:	5'- TTCGCCACTCATCCGATGTG
10	SEQ ID No. 221:	5'- CATTTAACGGGATGCGTTCG
	SEQ ID No. 222:	5'- CACAGTTCGCCACTCATCCG
	SEQ ID No. 223:	5'- TTCGCCACAGTTCGCCACTC
	SEQ ID No. 224:	5'- CTCCGCGTTTGTACACCGGCA
	SEQ ID No. 225:	5'- ACGCCGCCAGGGTTCATCCT
15	SEQ ID No. 226:	5'- CCTTCCAGAAGTGATAGCCG
	SEQ ID No. 227:	5'- TCATTGCCTTGGTGAACCGT
	SEQ ID No. 228:	5'- CACAGTATGTCAAGACCTGG
	SEQ ID No. 229:	5'- TTGGTGAACCGTTACTCCAC
	SEQ ID No. 230:	5'- CTTGGTGAACCGTTACTCCA
20	SEQ ID No. 231:	5'- GTGAACCGTTACTCCACCAA
	SEQ ID No. 232:	5'- GGCTCCCGAAGGTTACCGAA
	SEQ ID No. 233:	5'- GAAGGTTACCGAACCGAGTT
	SEQ ID No. 234:	5'- TGGCTCCCGAAGGTTACCGA
	SEQ ID No. 235:	5'- TAATACGCCGCGGGTCCTTC
25	SEQ ID No. 236:	5'- GAACCGTTACTCCACCAACT
	SEQ ID No. 237:	5'- TACGCCGCGGGTCCTTCCAG
	SEQ ID No. 238:	5'- TCACCAGTTCGCCACAGTTC
	SEQ ID No. 239:	5'- CCTTGGTGAACCGTTACTCC
	SEQ ID No. 240:	5'- CTCACCAGTTCGCCACAGTT
30	SEQ ID No. 241:	5'- CGCCGCCAGGGTTCATCCTG



SEQ ID No. 242: 5'- CCTTGGTGAACCATTA CTCC  
SEQ ID No. 243: 5'- TGGTGAACCATTA CTCCACC  
SEQ ID No. 244: 5'- GCCGCCAGGGTTCATCCTGA  
SEQ ID No. 245: 5'- GGTGAACCATTA CTCCACCA  
5 SEQ ID No. 246: 5'- CCAGGGTTCATCCTGAGCCA  
SEQ ID No. 247: 5'- AATACGCCGCGGGTCCTTCC  
SEQ ID No. 248: 5'- CACGCCGCCAGGGTTCATCC  
SEQ ID No. 249: 5'- AGTTCGCCACTCATCCGATG  
SEQ ID No. 250: 5'- CGGGATGCGTTCGACTTGCA  
10 SEQ ID No. 251: 5'- CATTGCCTTGGTGAACCGTT  
SEQ ID No. 252: 5'- GCACGCCGCCAGGGTTCATC  
SEQ ID No. 253: 5'- CTTCTCCGCGTTTGTCACC  
SEQ ID No. 254: 5'- TGGTGAACCGTTACTCCACC  
SEQ ID No. 255: 5'- CCTTCCTCCGCGTTTGTCAC  
15 SEQ ID No. 256: 5'- ACGCCGCGGGTCCTTCCAGA  
SEQ ID No. 257: 5'- GGTGAACCGTTACTCCACCA  
SEQ ID No. 258: 5'- GGGTCCTTCCAGAAGTGATA  
SEQ ID No. 259: 5'- CTTCCAGAAGTGATAGCCGA  
SEQ ID No. 260: 5'- GCCTTGGTGAACCATTA CT  
20 SEQ ID No. 261: 5'- ACAGTTCGCCACTCATCCGA  
SEQ ID No. 262: 5'- ACCTTCCTCCGCGTTTGTCAC  
SEQ ID No. 263: 5'- CGAACCGACTTTGGGTGTTG  
SEQ ID No. 264: 5'- GAACCGACTTTGGGTGTTGC  
SEQ ID No. 265: 5'- AGGTTACCGAACCGACTTTG  
25 SEQ ID No. 266: 5'- ACCGAACCGACTTTGGGTGT  
SEQ ID No. 267: 5'- TTACCGAACCGACTTTGGGT  
SEQ ID No. 268: 5'- TACCGAACCGACTTTGGGTG  
SEQ ID No. 269: 5'- GTTACCGAACCGACTTTGGG  
SEQ ID No. 270: 5'- CCTTTCTGGTATGGTACCGTC  
30 SEQ ID No. 271: 5'- TGCACCGCGGAYCCATCTCT

SEQ ID No. 272: 5'- AGTTGCAGTCCAGTAAGCCG  
SEQ ID No. 273: 5'- GTTGCAGTCCAGTAAGCCGC  
SEQ ID No. 274: 5'- CAGTTGCAGTCCAGTAAGCC  
SEQ ID No. 275: 5'- TGCAGTCCAGTAAGCCGCCT  
5 SEQ ID No. 276: 5'- TCAGTTGCAGTCCAGTAAGC  
SEQ ID No. 277: 5'- TTGCAGTCCAGTAAGCCGCC  
SEQ ID No. 278: 5'- GCAGTCCAGTAAGCCGCCTT  
SEQ ID No. 279: 5'- GTCAGTTGCAGTCCAGTAAG  
SEQ ID No. 280: 5'- CTCTAGGTGACGCCGAAGCG  
10 SEQ ID No. 281: 5'- ATCTCTAGGTGACGCCGAAG  
SEQ ID No. 282: 5'- TCTAGGTGACGCCGAAGCGC  
SEQ ID No. 283: 5'- TCTCTAGGTGACGCCGAAGC  
SEQ ID No. 284: 5'- CCATCTCTAGGTGACGCCGA  
SEQ ID No. 285: 5'- CATCTCTAGGTGACGCCGAA  
15 SEQ ID No. 286: 5'- TAGGTGACGCCGAAGCGCCT  
SEQ ID No. 287: 5'- CTAGGTGACGCCGAAGCGCC  
SEQ ID No. 288: 5'- CTTAGACGGCTCCTTCCTAA  
SEQ ID No. 289: 5'- CCTTAGACGGCTCCTTCCTA  
SEQ ID No. 290: 5'- ACGTCAGTTGCAGTCCAGTA  
20 SEQ ID No. 291: 5'- CGTCAGTTGCAGTCCAGTAA  
SEQ ID No. 292: 5'- ACGCCGAAGCGCCTTTTAAC  
SEQ ID No. 293: 5'- GACGCCGAAGCGCCTTTTAA  
SEQ ID No. 294: 5'- GCCGAAGCGCCTTTTAACTT  
SEQ ID No. 295: 5'- CGCCGAAGCGCCTTTTAACT  
25 SEQ ID No. 296: 5'- GTGACGCCGAAGCGCCTTTT  
SEQ ID No. 297: 5'- TGACGCCGAAGCGCCTTTTA  
SEQ ID No. 298: 5'- AGACGGCTCCTTCCTAAAAG  
SEQ ID No. 299: 5'- ACGGCTCCTTCCTAAAAGGT  
SEQ ID No. 300: 5'- GACGGCTCCTTCCTAAAAGG  
30 SEQ ID No. 301: 5'- CCTTCCTAAAAGGTTAGGCC

	SEQ ID No. 302:	5'- GGTGACGCCAAAGCGCCTTT
	SEQ ID No. 303:	5'- AGGTGACGCCAAAGCGCCTT
	SEQ ID No. 304:	5'- TAGGTGACGCCAAAGCGCCT
	SEQ ID No. 305:	5'- CTCTAGGTGACGCCAAAGCG
5	SEQ ID No. 306:	5'- TCTAGGTGACGCCAAAGCGC
	SEQ ID No. 307:	5'- CTAGGTGACGCCAAAGCGCC
	SEQ ID No. 308:	5'- ACGCCAAAGCGCCTTTTAAC
	SEQ ID No. 309:	5'- CGCCAAAGCGCCTTTTAACT
	SEQ ID No. 310:	5'- TGACGCCAAAGCGCCTTTTA
10	SEQ ID No. 311:	5'- TCTCTAGGTGACGCCAAAGC
	SEQ ID No. 312:	5'- GTGACGCCAAAGCGCCTTTT
	SEQ ID No. 313:	5'- GACGCCAAAGCGCCTTTTAA
	SEQ ID No. 314:	5'- ATCTCTAGGTGACGCCAAAG
	SEQ ID No. 315:	5'- CATCTCTAGGTGACGCCAAA
15	SEQ ID No. 316:	5'- TCCATCTCTAGGTGACGCCA
	SEQ ID No. 317:	5'- CCATCTCTAGGTGACGCCAA
	SEQ ID No. 318:	5'- CTGCCTTAGACGGCTCCCCC
	SEQ ID No. 319:	5'- CCTGCCTTAGACGGCTCCCC
	SEQ ID No. 320:	5'- GTGTCATGCGACACTGAGTT
20	SEQ ID No. 321:	5'- TGTGTCATGCGACACTGAGT
	SEQ ID No. 322:	5'- CTTTGTGTCATGCGACACTG
	SEQ ID No. 323:	5'- TTGTGTCATGCGACACTGAG
	SEQ ID No. 324:	5'- TGCCTTAGACGGCTCCCCCT
	SEQ ID No. 325:	5'- AGACGGCTCCCCCTAAAAGG
25	SEQ ID No. 326:	5'- TAGACGGCTCCCCCTAAAAG
	SEQ ID No. 327:	5'- GCCTTAGACGGCTCCCCCTA
	SEQ ID No. 328:	5'- GCTCCCCCTAAAAGGTTAGG
	SEQ ID No. 329:	5'- GGCTCCCCCTAAAAGGTTAG
	SEQ ID No. 330:	5'- CTCCCCCTAAAAGGTTAGGC
30	SEQ ID No. 331:	5'- TCCCCCTAAAAGGTTAGGCC

SEQ ID No. 332: 5'- CCCTAAAAGGTTAGGCCACC  
SEQ ID No. 333: 5'- CCCCTAAAAGGTTAGGCCAC  
SEQ ID No. 334: 5'- CGGCTCCCCCTAAAAGGTTA  
SEQ ID No. 335: 5'- CCCCCTAAAAGGTTAGGCCA  
5 SEQ ID No. 336: 5'- CTTAGACGGCTCCCCCTAAA  
SEQ ID No. 337: 5'- TTAGACGGCTCCCCCTAAAA  
SEQ ID No. 338: 5'- GGGTTCGCAACTCGTTGTAT  
SEQ ID No. 339: 5'- CCTTAGACGGCTCCCCCTAA  
SEQ ID No. 340: 5'- ACGGCTCCCCCTAAAAGGTT  
10 SEQ ID No. 341: 5'- GACGGCTCCCCCTAAAAGGT  
SEQ ID No. 342: 5'- ACGCCGCAAGACCATCCTCT  
SEQ ID No. 343: 5'- CTAATACGCCGCAAGACCAT  
SEQ ID No. 344: 5'- TACGCCGCAAGACCATCCTC  
SEQ ID No. 345: 5'- GTTACGATCTAGCAAGCCGC  
15 SEQ ID No. 346: 5'- AATACGCCGCAAGACCATCC  
SEQ ID No. 347: 5'- CGCCGCAAGACCATCCTCTA  
SEQ ID No. 348: 5'- GCTAATACGCCGCAAGACCA  
SEQ ID No. 349: 5'- ACCATCCTCTAGCGATCCAA  
SEQ ID No. 350: 5'- TAATACGCCGCAAGACCATC  
20 SEQ ID No. 351: 5'- AGCCATCCCTTTCTGGTAAG  
SEQ ID No. 352: 5'- ATACGCCGCAAGACCATCCT  
SEQ ID No. 353: 5'- AGTTACGATCTAGCAAGCCG  
SEQ ID No. 354: 5'- AGCTAATACGCCGCAAGACC  
SEQ ID No. 355: 5'- GCCGCAAGACCATCCTCTAG  
25 SEQ ID No. 356: 5'- TTACGATCTAGCAAGCCGCT  
SEQ ID No. 357: 5'- GACCATCCTCTAGCGATCCA  
SEQ ID No. 358: 5'- TTGCTACGTCACTAGGAGGC  
SEQ ID No. 359: 5'- ACGTCACTAGGAGGCGGAAA  
SEQ ID No. 360: 5'- TTTGCTACGTCACTAGGAGG  
30 SEQ ID No. 361: 5'- GCCATCCCTTTCTGGTAAGG

SEQ ID No. 362: 5'- TACGTCACTAGGAGGCGGAA  
SEQ ID No. 363: 5'- CGTCACTAGGAGGCGGAAAC  
SEQ ID No. 364: 5'- AAGACCATCCTCTAGCGATC  
SEQ ID No. 365: 5'- GCACGTATTTAGCCATCCCT  
5 SEQ ID No. 366: 5'- CTCTAGCGATCCAAAAGGAC  
SEQ ID No. 367: 5'- CCTCTAGCGATCCAAAAGGA  
SEQ ID No. 368: 5'- CCATCCTCTAGCGATCCAAA  
SEQ ID No. 369: 5'- GGCACGTATTTAGCCATCCC  
SEQ ID No. 370: 5'- TACGATCTAGCAAGCCGCTT  
10 SEQ ID No. 371: 5'- CAGTTACGATCTAGCAAGCC  
SEQ ID No. 372: 5'- CCGCAAGACCATCCTCTAGC  
SEQ ID No. 373: 5'- CCATCCCTTTCTGGTAAGGT  
SEQ ID No. 374: 5'- AGACCATCCTCTAGCGATCC  
SEQ ID No. 375: 5'- CAAGACCATCCTCTAGCGAT  
15 SEQ ID No. 376: 5'- GCTACGTCACTAGGAGGCGG  
SEQ ID No. 377: 5'- TGCTACGTCACTAGGAGGCG  
SEQ ID No. 378: 5'- CTACGTCACTAGGAGGCGGA  
SEQ ID No. 379: 5'- CCTCAACGTCAGTTACGATC  
SEQ ID No. 380: 5'- GTCACTAGGAGGCGGAAACC  
20 SEQ ID No. 381: 5'- TCCTCTAGCGATCCAAAAGG  
SEQ ID No. 382: 5'- TGGCACGTATTTAGCCATCC  
SEQ ID No. 383: 5'- ACGATCTAGCAAGCCGCTTT  
SEQ ID No. 384: 5'- GCCAGTCTCTCAACTCGGCT  
SEQ ID No. 385: 5'- AAGCTAATACGCCGCAAGAC  
25 SEQ ID No. 386: 5'- GTTTGCTACGTCACTAGGAG  
SEQ ID No. 387: 5'- CGCCACTCTAGTCATTGCCT  
SEQ ID No. 388: 5'- GGCCAGCCAGTCTCTCAACT  
SEQ ID No. 389: 5'- CAGCCAGTCTCTCAACTCGG  
SEQ ID No. 390: 5'- CCCGAAGATCAATTCAGCGG  
30 SEQ ID No. 391: 5'- CCGGCCAGTCTCTCAACTCG

SEQ ID No. 392: 5'- CCAGCCAGTCTCTCAACTCG  
SEQ ID No. 393: 5'- TCATTGCCTCACTTCACCCG  
SEQ ID No. 394: 5'- GCCAGCCAGTCTCTCAACTC  
SEQ ID No. 395: 5'- CACCCGAAGATCAATTCAGC  
5 SEQ ID No. 396: 5'- GTCATTGCCTCACTTCACCC  
SEQ ID No. 397: 5'- CATTGCCTCACTTCACCCGA  
SEQ ID No. 398: 5'- ATTGCCTCACTTCACCCGAA  
SEQ ID No. 399: 5'- CGAAGATCAATTCAGCGGCT  
SEQ ID No. 400: 5'- AGTCATTGCCTCACTTCACC  
10 SEQ ID No. 401: 5'- TCGCCACTCTAGTCATTGCC  
SEQ ID No. 402: 5'- TTGCCTCACTTCACCCGAAG  
SEQ ID No. 403: 5'- CGGCCAGTCTCTCAACTCGG  
SEQ ID No. 404: 5'- CTGGCACGTATTTAGCCATC  
SEQ ID No. 405: 5'- ACCCGAAGATCAATTCAGCG  
15 SEQ ID No. 406: 5'- TCTAGCGATCCAAAAGGACC  
SEQ ID No. 407: 5'- CTAGCGATCCAAAAGGACCT  
SEQ ID No. 408: 5'- GCACCCATCGTTTACGGTAT  
SEQ ID No. 409: 5'- CACCCATCGTTTACGGTATG  
SEQ ID No. 410: 5'- GCCACTCTAGTCATTGCCTC  
20 SEQ ID No. 411: 5'- CGTTTGCTACGTCCTAGGA  
SEQ ID No. 412: 5'- GCCTCAACGTCAGTTACGAT  
SEQ ID No. 413: 5'- GCCGGCCAGTCTCTCAACTC  
SEQ ID No. 414: 5'- TCACTAGGAGGCGGAAACCT  
SEQ ID No. 415: 5'- AGCCTCAACGTCAGTTACGA  
25 SEQ ID No. 416: 5'- AGCCAGTCTCTCAACTCGGC  
SEQ ID No. 417: 5'- GGCCAGTCTCTCAACTCGGC  
SEQ ID No. 418: 5'- CAAGCTAATACGCCGCAAGA  
SEQ ID No. 419: 5'- TTCGCCACTCTAGTCATTGC  
SEQ ID No. 420: 5'- CCGAAGATCAATTCAGCGGC  
30 SEQ ID No. 421: 5'- CGCAAGACCATCCTCTAGCG

SEQ ID No. 422: 5'- GCAAGACCATCCTCTAGCGA  
SEQ ID No. 423: 5'- GCGTTTGCTACGTCACTAGG  
SEQ ID No. 424: 5'- CCACTCTAGTCATTGCCTCA  
SEQ ID No. 425: 5'- CACTCTAGTCATTGCCTCAC  
5 SEQ ID No. 426: 5'- CCAGTCTCTCAACTCGGCTA  
SEQ ID No. 427: 5'- TTACCTTAGGCACCGGCCTC  
SEQ ID No. 428: 5'- ACAAGCTAATACGCCGCAAG  
SEQ ID No. 429: 5'- TTTACCTTAGGCACCGGCCT  
SEQ ID No. 430: 5'- TTTTACCTTAGGCACCGGCC  
10 SEQ ID No. 431: 5'- ATTTTACCTTAGGCACCGGC  
SEQ ID No. 432: 5'- GATTTTACCTTAGGCACCGG  
SEQ ID No. 433: 5'- CTCACTTCACCCGAAGATCA  
SEQ ID No. 434: 5'- ACGCCACCAGCGTTCATCCT  
SEQ ID No. 435: 5'- GCCAAGCGACTTTGGGTACT  
15 SEQ ID No. 436: 5'- CGGAAAATTCCCTACTGCAG  
SEQ ID No. 437: 5'- CGATCTAGCAAGCCGCTTTC  
SEQ ID No. 438: 5'- GGTACCGTCAAGCTGAAAAC  
SEQ ID No. 439: 5'- TGCCTCACTTCACCCGAAGA  
SEQ ID No. 440: 5'- GGCCGGCCAGTCTCTCAACT  
20 SEQ ID No. 441: 5'- GGTAAGGTACCGTCAAGCTG  
SEQ ID No. 442: 5'- GTAAGGTACCGTCAAGCTGA  
SEQ ID No. 443: 5'- CCGCAAGACCATCCTCTAGG  
SEQ ID No. 444: 5'- ATTTAGCCATCCCTTTCTGG  
SEQ ID No. 445: 5'- AACCCTTCATCACACACG  
25 SEQ ID No. 446: 5'- CGAAACCCTTCATCACAC  
SEQ ID No. 447: 5'- ACCCTTCATCACACACGC  
SEQ ID No. 448: 5'- TACCGTCACACACTGAAC  
SEQ ID No. 449: 5'- AGATACCGTCACACACTG  
SEQ ID No. 450: 5'- CACTCAAGGGCGGAAACC  
30 SEQ ID No. 451: 5'- ACCGTCACACACTGAACA

SEQ ID No. 452: 5'- CGTCACACACTGAACAGT  
SEQ ID No. 453: 5'- CCGAAACCCTTCATCACA  
SEQ ID No. 454: 5'- CCGTCACACACTGAACAG  
SEQ ID No. 455: 5'- GATACCGTCACACACTGA  
5 SEQ ID No. 456: 5'- GGTAAGATACCGTCACAC  
SEQ ID No. 457: 5'- CCCTTCATCACACACGCG  
SEQ ID No. 458: 5'- ACAGTGTTTTACGAGCCG  
SEQ ID No. 459: 5'- CAGTGTTTTACGAGCCGA  
SEQ ID No. 460: 5'- ACAAAGCGTTCGACTTGC  
10 SEQ ID No. 461: 5'- CGGATAACGCTTGGAACA  
SEQ ID No. 462: 5'- AGGGCGGAAACCCTCGAA  
SEQ ID No. 463: 5'- GGGCGGAAACCCTCGAAC  
SEQ ID No. 464: 5'- GGC GGAAACCCTCGAACA  
SEQ ID No. 465: 5'- TGAGGGCTTTCACTTCAG  
15 SEQ ID No. 466: 5'- AGGGCTTTCACTTCAGAC  
SEQ ID No. 467: 5'- GAGGGCTTTCACTTCAGA  
SEQ ID No. 468: 5'- ACTGCACTCAAGTCATCC  
SEQ ID No. 469: 5'- CCGGATAACGCTTGGAAC  
SEQ ID No. 470: 5'- TCCGGATAACGCTTGGA  
20 SEQ ID No. 471: 5'- TATCCCCTGCTAAGAGGT  
SEQ ID No. 472: 5'- CCTGCTAAGAGGTAGGT  
SEQ ID No. 473: 5'- CCCTGCTAAGAGGTAGGT  
SEQ ID No. 474: 5'- CCCCTGCTAAGAGGTAGG  
SEQ ID No. 475: 5'- TCCCCTGCTAAGAGGTAG  
25 SEQ ID No. 476: 5'- ATCCCCTGCTAAGAGGTA  
SEQ ID No. 477: 5'- CCGTTCCTTTCTGGTAAG  
SEQ ID No. 478: 5'- GCCGTTCCTTTCTGGTAA  
SEQ ID No. 479: 5'- AGCCGTTCCTTTCTGGTA  
SEQ ID No. 480: 5'- GCACGTATTTAGCCGTTC  
30 SEQ ID No. 481: 5'- CACGTATTTAGCCGTTC



SEQ ID No. 482: 5'-GGCACGTATTTAGCCGTT  
SEQ ID No. 483: 5'-CACTTTCCTCTACTGCAC  
SEQ ID No. 484: 5'-CCACTTTCCTCTACTGCA  
SEQ ID No. 485: 5'-TCCACTTTCCTCTACTGC  
5 SEQ ID No. 486: 5'-CTTTCCTCTACTGCACTC  
SEQ ID No. 487: 5'-TAGCCGTTTCCTTTCTGGT  
SEQ ID No. 488: 5'-TTAGCCGTTTCCTTTCTGG  
SEQ ID No. 489: 5'-TTATCCCCTGCTAAGAGG  
SEQ ID No. 490: 5'-GTTATCCCCTGCTAAGAG  
10 SEQ ID No. 491: 5'-CCCGTTTCGCCACTCTTTG  
SEQ ID No. 492: 5'-AGCTGAGGGCTTTCACTT  
SEQ ID No. 493: 5'-GAGCTGAGGGCTTTCACT  
SEQ ID No. 494: 5'-GCTGAGGGCTTTCACTTC  
SEQ ID No. 495: 5'-CTGAGGGCTTTCACTTCA  
15 SEQ ID No. 496: 5'-CCCGTGTCCCGAAGGAAC  
SEQ ID No. 497: 5'-GCACGAGTATGTCAAGAC  
SEQ ID No. 498: 5'-GTATCCCGTGTCCCGAAG  
SEQ ID No. 499: 5'-TCCCGTGTCCCGAAGGAA  
SEQ ID No. 500: 5'-ATCCCGTGTCCCGAAGGA  
20 SEQ ID No. 501: 5'-TATCCCGTGTCCCGAAGG  
SEQ ID No. 502: 5'-CTTACCTTAGGAAGCGCC  
SEQ ID No. 503: 5'-TTACCTTAGGAAGCGCCC  
SEQ ID No. 504: 5'-CCTGTATCCCGTGTCCCG  
SEQ ID No. 505: 5'-CCACCTGTATCCCGTGTCC  
25 SEQ ID No. 506: 5'-CACCTGTATCCCGTGTCC  
SEQ ID No. 507: 5'-ACCTGTATCCCGTGTCCC  
SEQ ID No. 508: 5'-CTGTATCCCGTGTCCCGA  
SEQ ID No. 509: 5'-TGTATCCCGTGTCCCGAA  
SEQ ID No. 510: 5'-CACGAGTATGTCAAGACC  
30 SEQ ID No. 511: 5'-CGGTCTTACCTTAGGAAG

SEQ ID No. 512: 5' TAGGAAGCGCCCTCCTTG  
SEQ ID No. 513: 5' AGGAAGCGCCCTCCTTGC  
SEQ ID No. 514: 5' TTAGGAAGCGCCCTCCTT  
SEQ ID No. 515: 5' CTTAGGAAGCGCCCTCCT  
5 SEQ ID No. 516: 5' CCTTAGGAAGCGCCCTCC  
SEQ ID No. 517: 5' ACCTTAGGAAGCGCCCTC  
SEQ ID No. 518: 5' TGCACACAATGGTTGAGC  
SEQ ID No. 519: 5' TACCTTAGGAAGCGCCCT  
SEQ ID No. 520: 5' ACCACCTGTATCCCGTGT  
10 SEQ ID No. 521: 5' GCACCACCTGTATCCCGT  
SEQ ID No. 522: 5' CACCACCTGTATCCCGTG  
SEQ ID No. 523: 5' GCGGTTAGGCAACCTACT  
SEQ ID No. 524: 5' TGCGGTTAGGCAACCTAC  
SEQ ID No. 525: 5' TTGCGGTTAGGCAACCTA  
15 SEQ ID No. 526: 5' GGTCTTACCTTAGGAAGC  
SEQ ID No. 527: 5' GCTAATACAACGCGGGAT  
SEQ ID No. 528: 5' CTAATACAACGCGGGATC  
SEQ ID No. 529: 5' ATACAACGCGGGATCATC  
SEQ ID No. 530: 5' CGGTTAGGCAACCTACTT  
20 SEQ ID No. 531: 5' TGCACCACCTGTATCCCG  
SEQ ID No. 532: 5' GAAGCGCCCTCCTTGCGG  
SEQ ID No. 533: 5' GGAAGCGCCCTCCTTGCG  
SEQ ID No. 534: 5' CGTCCCTTTCTGGTTAGA  
SEQ ID No. 535: 5' AGCTAATACAACGCGGGA  
25 SEQ ID No. 536: 5' TAGCTAATACAACGCGGG  
SEQ ID No. 537: 5' CTAGCTAATACAACGCGG  
SEQ ID No. 538: 5' GGCTATGTATCATCGCCT  
SEQ ID No. 539: 5' GAGCCACTGCCTTTTACA  
SEQ ID No. 540: 5' GTCGGCTATGTATCATCG  
30 SEQ ID No. 541: 5' GGTCGGCTATGTATCATC

SEQ ID No. 542: 5' CAGGTCGGCTATGTATCA  
SEQ ID No. 543: 5' CGGCTATGTATCATCGCC  
SEQ ID No. 544: 5' TCGGCTATGTATCATCGC  
SEQ ID No. 545: 5' GTCTTACCTTAGGAAGCG  
5 SEQ ID No. 546: 5' TCTTACCTTAGGAAGCGC  
SEQ ID No. 547: 5'- GTACAAACCGCCTACACGCC  
SEQ ID No. 548: 5'- TGTACAAACCGCCTACACGC  
SEQ ID No. 549: 5'- GATCAGCACGATGTCGCCAT  
SEQ ID No. 550: 5'- CTGTACAAACCGCCTACACG  
10 SEQ ID No. 551: 5'- GAGATCAGCACGATGTCGCC  
SEQ ID No. 552: 5'- AGATCAGCACGATGTCGCCA  
SEQ ID No. 553: 5'- ATCAGCACGATGTCGCCATC  
SEQ ID No. 554: 5'- TCAGCACGATGTCGCCATCT  
SEQ ID No. 555: 5'- ACTGTACAAACCGCCTACAC  
15 SEQ ID No. 556: 5'- CCGCCACTAAGGCCGAAACC  
SEQ ID No. 557: 5'- CAGCACGATGTCGCCATCTA  
SEQ ID No. 558: 5'- TACAAACCGCCTACACGCCC  
SEQ ID No. 559: 5'- AGCACGATGTCGCCATCTAG  
SEQ ID No. 560: 5'- CGGCTTTTAGAGATCAGCAC  
20 SEQ ID No. 561: 5'- TCCGCCACTAAGGCCGAAAC  
SEQ ID No. 562: 5'- GACTGTACAAACCGCCTACA  
SEQ ID No. 563: 5'- GTCCGCCACTAAGGCCGAAA  
SEQ ID No. 564: 5'- GGGGATTTACATCTGACTG  
SEQ ID No. 565: 5'- CATACAAGCCCTGGTAAGGT  
25 SEQ ID No. 566: 5'- ACAAGCCCTGGTAAGGTTCT  
SEQ ID No. 567: 5'- ACAAACCGCCTACACGCCCT  
SEQ ID No. 568: 5'- CTGACTGTACAAACCGCCTA  
SEQ ID No. 569: 5'- TGACTGTACAAACCGCCTAC  
SEQ ID No. 570: 5'- ACGATGTCGCCATCTAGCTT  
30 SEQ ID No. 571: 5'- CACGATGTCGCCATCTAGCT

SEQ ID No. 572: 5'-CGATGTCGCCATCTAGCTTC  
SEQ ID No. 573: 5'-GCACGATGTCGCCATCTAGC  
SEQ ID No. 574: 5'-GATGTCGCCATCTAGCTTCC  
SEQ ID No. 575: 5'-ATGTCGCCATCTAGCTTCCC  
5 SEQ ID No. 576: 5'-TGTCGCCATCTAGCTTCCCA  
SEQ ID No. 577: 5'-GCCATCTAGCTTCCCCTGT  
SEQ ID No. 578: 5'-TCGCCATCTAGCTTCCCCT  
SEQ ID No. 579: 5'-CGCCATCTAGCTTCCCCTG  
SEQ ID No. 580: 5'-GTCGCCATCTAGCTTCCCAC  
10 SEQ ID No. 581: 5'-TACAAGCCCTGGTAAGGTTC  
SEQ ID No. 582: 5'-GCCACTAAGGCCGAAACCTT  
SEQ ID No. 583: 5'-ACTAAGGCCGAAACCTTCGT  
SEQ ID No. 584: 5'-CTAAGGCCGAAACCTTCGTG  
SEQ ID No. 585: 5'-CACTAAGGCCGAAACCTTCG  
15 SEQ ID No. 586: 5'-AAGGCCGAAACCTTCGTGCG  
SEQ ID No. 587: 5'-CCACTAAGGCCGAAACCTTC  
SEQ ID No. 588: 5'-TAAGGCCGAAACCTTCGTGC  
SEQ ID No. 589: 5'-AGGCCGAAACCTTCGTGCGA  
SEQ ID No. 590: 5'-TCTGACTGTACAAACCGCCT  
20 SEQ ID No. 591: 5'-CATCTGACTGTACAAACCGC  
SEQ ID No. 592: 5'-ATCTGACTGTACAAACCGCC  
SEQ ID No. 593: 5'-CTTCGTGCGACTTGCGATGTG  
SEQ ID No. 594: 5'-CCTTCGTGCGACTTGCGATGT  
SEQ ID No. 595: 5'-CTCTCTAGAGTGCCCACCCA  
25 SEQ ID No. 596: 5'-TCTCTAGAGTGCCCACCCAA  
SEQ ID No. 597: 5'-ACGTATCAAATGCAGCTCCC  
SEQ ID No. 598: 5'-CGTATCAAATGCAGCTCCCA  
SEQ ID No. 599: 5'-CGCCACTAAGGCCGAAACCT  
SEQ ID No. 600: 5'-CCGAAACCTTCGTGCGACTT  
30 SEQ ID No. 601: 5'-GCCGAAACCTTCGTGCGACT

SEQ ID No. 602: 5'- AACCTTCGTGCGACTTG CAT  
SEQ ID No. 603: 5'- CGAAACCTTCGTGCGACTTG  
SEQ ID No. 604: 5'- ACCTTCGTGCGACTTG CATG  
SEQ ID No. 605: 5'- GAAACCTTCGTGCGACTTG C  
5 SEQ ID No. 606: 5'- GGCCGAAACCTTCGTGCGAC  
SEQ ID No. 607: 5'- AAACCTTCGTGCGACTTG CA  
SEQ ID No. 608: 5'- CACGTATCAAATGCAGCTCC  
SEQ ID No. 609: 5'- GCTCACCGGCTTAAGGTCAA  
SEQ ID No. 610: 5'- CGCTCACCGGCTTAAGGTCA  
10 SEQ ID No. 611: 5'- TCGCTCACCGGCTTAAGGTC  
SEQ ID No. 612: 5'- CTCACCGGCTTAAGGTCAAA  
SEQ ID No. 613: 5'- CCCGACCGTGGTCGGCTGCG  
SEQ ID No. 614: 5'- GCTCACCGGCTTAAGGTCAA  
SEQ ID No. 615: 5'- CGCTCACCGGCTTAAGGTCA  
15 SEQ ID No. 616: 5'- TCGCTCACCGGCTTAAGGTC  
SEQ ID No. 617: 5'- CTCACCGGCTTAAGGTCAAA  
SEQ ID No. 618: 5'- CCCGACCGTGGTCGGCTGCG  
SEQ ID No. 619: 5'- TCACCGGCTTAAGGTCAAAC  
SEQ ID No. 620: 5'- CAACCCTCTCTCACACTCTA  
20 SEQ ID No. 621: 5'- ACAACCCTCTCTCACACTCT  
SEQ ID No. 622: 5'- CCACAACCCTCTCTCACACT  
SEQ ID No. 623: 5'- AACCCTCTCTCACACTCTAG  
SEQ ID No. 624: 5'- CACAACCCTCTCTCACACTC  
SEQ ID No. 625: 5'- TCCACAACCCTCTCTCACAC  
25 SEQ ID No. 626: 5'- TTCCACAACCCTCTCTCACA  
SEQ ID No. 627: 5'- ACCCTCTCTCACACTCTAGT  
SEQ ID No. 628: 5'- GAGCCAGGTTGCCGCCTTCG  
SEQ ID No. 629: 5'- AGGTCAAACCAACTCCCATG  
SEQ ID No. 630: 5'- ATGAGCCAGGTTGCCGCCTT  
30 SEQ ID No. 631: 5'- TGAGCCAGGTTGCCGCCTTC

SEQ ID No. 632: 5'- AGGCTCCTCCACAGGCGACT  
SEQ ID No. 633: 5'- CAGGCTCCTCCACAGGCGAC  
SEQ ID No. 634: 5'- GCAGGCTCCTCCACAGGCGA  
SEQ ID No. 635: 5'- TTCGCTCACCGGCTTAAGGT  
5 SEQ ID No. 636: 5'- GTTCGCTCACCGGCTTAAGG  
SEQ ID No. 637: 5'- GGTTCGCTCACCGGCTTAAG  
SEQ ID No. 638: 5'- ATTCCACAACCCTCTCTCAC  
SEQ ID No. 639: 5'- TGACCCGACCGTGGTCGGCT  
SEQ ID No. 640: 5'- CCCTCTCTCACACTCTAGTC  
10 SEQ ID No. 641: 5'- GAATTCCACAACCCTCTCTC  
SEQ ID No. 642: 5'- AGCCAGGTTGCCGCCTTCGC  
SEQ ID No. 643: 5'- GCCAGGTTGCCGCCTTCGCC  
SEQ ID No. 644: 5'- GGAATTCCACAACCCTCTCT  
SEQ ID No. 645: 5'- GGGAATTCCACAACCCTCTC  
15 SEQ ID No. 646: 5'- AACGCAGGCTCCTCCACAGG  
SEQ ID No. 647: 5'- CGGCTTAAGGTCAAACCAAC  
SEQ ID No. 648: 5'- CCGGCTTAAGGTCAAACCAA  
SEQ ID No. 649: 5'- CACCGGCTTAAGGTCAAACC  
SEQ ID No. 650: 5'- ACCGGCTTAAGGTCAAACCA  
20 SEQ ID No. 651: 5'- ACCCAACATCCAGCACACAT  
SEQ ID No. 652: 5'- TCGCTGACCCGACCGTGGTC  
SEQ ID No. 653: 5'- CGCTGACCCGACCGTGGTCG  
SEQ ID No. 654: 5'- GACCCGACCGTGGTCGGCTG  
SEQ ID No. 655: 5'- GCTGACCCGACCGTGGTCGG  
25 SEQ ID No. 656: 5'- CTGACCCGACCGTGGTCGGC  
SEQ ID No. 657: 5'- CAGGCGACTTGCGCCTTTGA  
SEQ ID No. 658: 5'- TCATGCGGTATTAGCTCCAG  
SEQ ID No. 659: 5'- ACTAGCTAATCGAACGCAGG  
SEQ ID No. 660: 5'- CATGCGGTATTAGCTCCAGT  
30 SEQ ID No. 661: 5'- CGCAGGCTCCTCCACAGGCG

SEQ ID No. 662: 5'- ACGCAGGCTCCTCCACAGGC  
SEQ ID No. 663: 5'- CTCAGGTGTCATGCGGTATT  
SEQ ID No. 664: 5'- CGCCTTTGACCCTCAGGTGT  
SEQ ID No. 665: 5'- ACCCTCAGGTGTCATGCGGT  
5 SEQ ID No. 666: 5'- CCTCAGGTGTCATGCGGTAT  
SEQ ID No. 667: 5'- TTTGACCCTCAGGTGTCATG  
SEQ ID No. 668: 5'- GACCCTCAGGTGTCATGCGG  
SEQ ID No. 669: 5'- TGACCCTCAGGTGTCATGCG  
SEQ ID No. 670: 5'- GCCTTTGACCCTCAGGTGTC  
10 SEQ ID No. 671: 5'- TTGACCCTCAGGTGTCATGC  
SEQ ID No. 672: 5'- CCCTCAGGTGTCATGCGGTA  
SEQ ID No. 673: 5'- CCTTTGACCCTCAGGTGTCA  
SEQ ID No. 674: 5'- CTTTGACCCTCAGGTGTCAT  
SEQ ID No. 675: 5'- AGTTATCCCCCACCCTATGGA  
15 SEQ ID No. 676: 5'- CCAGCTATCGATCATCGCCT  
SEQ ID No. 677: 5'- ACCAGCTATCGATCATCGCC  
SEQ ID No. 678: 5'- CAGCTATCGATCATCGCCTT  
SEQ ID No. 679: 5'- AGCTATCGATCATCGCCTTG  
SEQ ID No. 680: 5'- GCTATCGATCATCGCCTTGG  
20 SEQ ID No. 681: 5'- CTATCGATCATCGCCTTGGT  
SEQ ID No. 682: 5'- TTCGTGCGACTTGTCATGTGT  
SEQ ID No. 683: 5'- TCGATCATCGCCTTGGTAGG  
SEQ ID No. 684: 5'- ATCGATCATCGCCTTGGTAG  
SEQ ID No. 685: 5'- CACAGGCGACTTGCGCCTTT  
25 SEQ ID No. 686: 5'- CCACAGGCGACTTGCGCCTT  
SEQ ID No. 687: 5'- TCCACAGGCGACTTGCGCCT  
SEQ ID No. 688: 5'- TCCTCCACAGGCGACTTGCG  
SEQ ID No. 689: 5'- CCTCCACAGGCGACTTGCGC  
SEQ ID No. 690: 5'- CTCCACAGGCGACTTGCGCC  
30 SEQ ID No. 691: 5'- ACAGGCGACTTGCGCCTTTG

	SEQ ID No. 692:	5'- GCTCACCGGCTTAAGGTCAA
	SEQ ID No. 693:	5'- CGCTCACCGGCTTAAGGTCA
	SEQ ID No. 694:	5'- TCGCTCACCGGCTTAAGGTC
	SEQ ID No. 695:	5'- CTCACCGGCTTAAGGTCAAA
5	SEQ ID No. 696:	5'- CCCGACCGTGGTCGGCTGCG
	SEQ ID No. 697:	5'- TCACCGGCTTAAGGTCAAAC
	SEQ ID No. 698:	5'- CAACCCTCTCTCACACTCTA
	SEQ ID No. 699:	5'- ACAACCCTCTCTCACACTCT
	SEQ ID No. 700:	5'- CCACAACCCTCTCTCACACT
10	SEQ ID No. 701:	5'- AACCCTCTCTCACACTCTAG
	SEQ ID No. 702:	5'- CACAACCCTCTCTCACACTC
	SEQ ID No. 703:	5'- TCCACAACCCTCTCTCACAC
	SEQ ID No. 704:	5'- TTCCACAACCCTCTCTCACA
	SEQ ID No. 705:	5'- ACCCTCTCTCACACTCTAGT
15	SEQ ID No. 706:	5'- GAGCCAGGTTGCCGCCTTCG
	SEQ ID No. 707:	5'- AGGTCAAACCAACTCCCATG
	SEQ ID No. 708:	5'- ATGAGCCAGGTTGCCGCCTT
	SEQ ID No. 709:	5'- TGAGCCAGGTTGCCGCCTTC
	SEQ ID No. 710:	5'- AGGCTCCTCCACAGGCGACT
20	SEQ ID No. 711:	5'- CAGGCTCCTCCACAGGCGAC
	SEQ ID No. 712:	5'- GCAGGCTCCTCCACAGGCGA
	SEQ ID No. 713:	5'- TTCGCTCACCGGCTTAAGGT
	SEQ ID No. 714:	5'- GTTCGCTCACCGGCTTAAGG
	SEQ ID No. 715:	5'- GGTTGCTCACCGGCTTAAG
25	SEQ ID No. 716:	5'- ATTCCACAACCCTCTCTCAC
	SEQ ID No. 717:	5'- TGACCCGACCGTGGTCGGCT
	SEQ ID No. 718:	5'- CCCTCTCTCACACTCTAGTC
	SEQ ID No. 719:	5'- GAATTCCACAACCCTCTCTC
	SEQ ID No. 720:	5'- AGCCAGGTTGCCGCCTTCGC
30	SEQ ID No. 721:	5'- GCCAGGTTGCCGCCTTCGCC



	SEQ ID No. 722:	5'- GGAATTCCACAACCCTCTCT
	SEQ ID No. 723:	5'- GGGAATTCCACAACCCTCTC
	SEQ ID No. 724:	5'- AACGCAGGCTCCTCCACAGG
	SEQ ID No. 725:	5'- CGGCTTAAGGTCAAACCAAC
5	SEQ ID No. 726:	5'- CCGGCTTAAGGTCAAACCAA
	SEQ ID No. 727:	5'- CACCGGCTTAAGGTCAAACC
	SEQ ID No. 728:	5'- ACCGGCTTAAGGTCAAACCA
	SEQ ID No. 729:	5'- ACCCAACATCCAGCACACAT
	SEQ ID No. 730:	5'- TCGCTGACCCGACCGTGGTC
10	SEQ ID No. 731:	5'- CGCTGACCCGACCGTGGTCG
	SEQ ID No. 732:	5'- GACCCGACCGTGGTCGGCTG
	SEQ ID No. 733:	5'- GCTGACCCGACCGTGGTCGG
	SEQ ID No. 734:	5'- CTGACCCGACCGTGGTCGGC
	SEQ ID No. 735:	5'- CAGGCGACTTGCGCCTTTGA
15	SEQ ID No. 736:	5'- TCATGCGGTATTAGCTCCAG
	SEQ ID No. 737:	5'- ACTAGCTAATCGAACGCAGG
	SEQ ID No. 738:	5'- CATGCGGTATTAGCTCCAGT
	SEQ ID No. 739:	5'- CGCAGGCTCCTCCACAGGCG
	SEQ ID No. 740:	5'- ACGCAGGCTCCTCCACAGGC
20	SEQ ID No. 741:	5'- CTCAGGTGTCATGCGGTATT
	SEQ ID No. 742:	5'- CGCCTTTGACCCTCAGGTGT
	SEQ ID No. 743:	5'- ACCCTCAGGTGTCATGCGGT
	SEQ ID No. 744:	5'- CCTCAGGTGTCATGCGGTAT
	SEQ ID No. 745:	5'- TTTGACCCTCAGGTGTCATG
25	SEQ ID No. 746:	5'- GACCCTCAGGTGTCATGCGG
	SEQ ID No. 747:	5'- TGACCCTCAGGTGTCATGCG
	SEQ ID No. 748:	5'- GCCTTTGACCCTCAGGTGTC
	SEQ ID No. 749:	5'- TTGACCCTCAGGTGTCATGC
	SEQ ID No. 750:	5'- CCCTCAGGTGTCATGCGGTA
30	SEQ ID No. 751:	5'- CCTTTGACCCTCAGGTGTCA

SEQ ID No. 752: 5'- CTTTGACCCTCAGGTGTCAT  
SEQ ID No. 753: 5'- AGTTATCCCCCACCCATGGA  
SEQ ID No. 754: 5'- CCAGCTATCGATCATCGCCT  
SEQ ID No. 755: 5'- ACCAGCTATCGATCATCGCC  
5 SEQ ID No. 756: 5'- CAGCTATCGATCATCGCCTT  
SEQ ID No. 757: 5'- AGCTATCGATCATCGCCTTG  
SEQ ID No. 758: 5'- GCTATCGATCATCGCCTTGG  
SEQ ID No. 759: 5'- CTATCGATCATCGCCTTGGT  
SEQ ID No. 760: 5'- TTCGTGCGACTTGCGATGTGT  
10 SEQ ID No. 761: 5'- TCGATCATCGCCTTGGTAGG  
SEQ ID No. 762: 5'- ATCGATCATCGCCTTGGTAG  
SEQ ID No. 763: 5'- CACAGGCGACTTGCGCCTTT  
SEQ ID No. 764: 5'- CCACAGGCGACTTGCGCCTT  
SEQ ID No. 765: 5'- TCCACAGGCGACTTGCGCCT  
15 SEQ ID No. 766: 5'- TCCTCCACAGGCGACTTGCG  
SEQ ID No. 767: 5'- CCTCCACAGGCGACTTGCGC  
SEQ ID No. 768: 5'- CTCCACAGGCGACTTGCGCC  
SEQ ID No. 769: 5'- ACAGGCGACTTGCGCCTTTG  
SEQ ID No. 770: 5'- TCACCGGCTTAAGGTCAAAC  
20 SEQ ID No. 771: 5'- CAACCCTCTCTCACACTCTA  
SEQ ID No. 772: 5'- ACAACCCTCTCTCACACTCT  
SEQ ID No. 773: 5'- CCACAACCCTCTCTGACACT  
SEQ ID No. 774: 5'- AACCCTCTCTCACACTCTAG  
SEQ ID No. 775: 5'- CACAACCCTCTCTCACACTC  
25 SEQ ID No. 776: 5'- TCCACAACCCTCTCTCACAC  
SEQ ID No. 777: 5'- TTCCACAACCCTCTCTCACA  
SEQ ID No. 778: 5'- ACCCTCTCTCACACTCTAGT  
SEQ ID No. 779: 5'- GAGCCAGGTTGCCGCCTTCG  
SEQ ID No. 780: 5'- AGGTCAAACCAACTCCCATG  
30 SEQ ID No. 781: 5'- ATGAGCCAGGTTGCCGCCTT

	SEQ ID No. 782:	5'-TGAGCCAGGTTGCCGCCTTC
	SEQ ID No. 783:	5'-AGGCTCCTCCACAGGCGACT
	SEQ ID No. 784:	5'-CAGGCTCCTCCACAGGCGAC
	SEQ ID No. 785:	5'-GCAGGCTCCTCCACAGGCGA
5	SEQ ID No. 786:	5'-TTCGCTCACCGGCTTAAGGT
	SEQ ID No. 787:	5'-GTTCGCTCACCGGCTTAAGG
	SEQ ID No. 788:	5'-GGTTCGCTCACCGGCTTAAG
	SEQ ID No. 789:	5'-ATTCCACAACCCTCTCTCAC
	SEQ ID No. 790:	5'-TGACCCGACCGTGGTCGGCT
10	SEQ ID No. 791:	5'-CCCTCTCTCACACTCTAGTC
	SEQ ID No. 792:	5'-GAATTCCACAACCCTCTCTC
	SEQ ID No. 793:	5'-AGCCAGGTTGCCGCCTTCGC
	SEQ ID No. 794:	5'-GCCAGGTTGCCGCCTTCGCC
	SEQ ID No. 795:	5'-GGAATTCCACAACCCTCTCT
15	SEQ ID No. 796:	5'-GGGAATTCCACAACCCTCTC
	SEQ ID No. 797:	5'-AACGCAGGCTCCTCCACAGG
	SEQ ID No. 798:	5'-CGGCTTAAGGTCAAACCAAC
	SEQ ID No. 799:	5'-CCGGCTTAAGGTCAAACCAA
	SEQ ID No. 800:	5'-CACCGGCTTAAGGTCAAACC
20	SEQ ID No. 801:	5'-ACCGGCTTAAGGTCAAACCA
	SEQ ID No. 802:	5'-ACCCAACATCCAGCACACAT
	SEQ ID No. 803:	5'-TCGCTGACCCGACCGTGGTC
	SEQ ID No. 804:	5'-CGCTGACCCGACCGTGGTCG
	SEQ ID No. 805:	5'-GACCCGACCGTGGTCGGCTG
25	SEQ ID No. 806:	5'-GCTGACCCGACCGTGGTCGG
	SEQ ID No. 807:	5'-CTGACCCGACCGTGGTCGGC
	SEQ ID No. 808:	5'-CAGGCGACTTGCGCCTTTGA
	SEQ ID No. 809:	5'-TCATGCGGTATTAGCTCCAG
	SEQ ID No. 810:	5'-ACTAGCTAATCGAACGCAGG
30	SEQ ID No. 811:	5'-CATGCGGTATTAGCTCCAGT

SEQ ID No. 812: 5'- CGCAGGCTCCTCCACAGGCG  
SEQ ID No. 813: 5'- ACGCAGGCTCCTCCACAGGC  
SEQ ID No. 814: 5'- CTCAGGTGTCATGCGGTATT  
SEQ ID No. 815: 5'- CGCCTTTGACCCTCAGGTGT  
5 SEQ ID No. 816: 5'- ACCCTCAGGTGTCATGCGGT  
SEQ ID No. 817: 5'- CCTCAGGTGTCATGCGGTAT  
SEQ ID No. 818: 5'- TTTGACCCTCAGGTGTCATG  
SEQ ID No. 819: 5'- GACCCTCAGGTGTCATGCGG  
SEQ ID No. 820: 5'- TGACCCTCAGGTGTCATGCG  
10 SEQ ID No. 821: 5'- GCCTTTGACCCTCAGGTGTC  
SEQ ID No. 822: 5'- TTGACCCTCAGGTGTCATGC  
SEQ ID No. 823: 5'- CCCTCAGGTGTCATGCGGTA  
SEQ ID No. 824: 5'- CCTTTGACCCTCAGGTGTCA  
SEQ ID No. 825: 5'- CTTTGACCCTCAGGTGTCAT  
15 SEQ ID No. 826: 5'- AGTTATCCCCCACCCTATGGA  
SEQ ID No. 827: 5'- CCAGCTATCGATCATCGCCT  
SEQ ID No. 828: 5'- ACCAGCTATCGATCATCGCC  
SEQ ID No. 829: 5'- CAGCTATCGATCATCGCCTT  
SEQ ID No. 830: 5'- AGCTATCGATCATCGCCTTG  
20 SEQ ID No. 831: 5'- GCTATCGATCATCGCCTTGG  
SEQ ID No. 832: 5'- CTATCGATCATCGCCTTGGT  
SEQ ID No. 833: 5'- TTCGTGCGACTTGCATGTGT  
SEQ ID No. 834: 5'- TCGATCATCGCCTTGGTAGG  
SEQ ID No. 835: 5'- ATCGATCATCGCCTTGGTAG  
25 SEQ ID No. 836: 5'- CACAGGCGACTTGCGCCTTT  
SEQ ID No. 837: 5'- CCACAGGCGACTTGCGCCTT  
SEQ ID No. 838: 5'- TCCACAGGCGACTTGCGCCT  
SEQ ID No. 839: 5'- TCCTCCACAGGCGACTTGCG  
SEQ ID No. 840: 5'- CCTCCACAGGCGACTTGCGC  
30 SEQ ID No. 841: 5'- CTCCACAGGCGACTTGCGCC

	SEQ ID No. 842:	5'- ACAGGCGACTTGCGCCTTTG
	SEQ ID No. 843:	5'- AGCCCCGGTTTCCCGGCGTT
	SEQ ID No. 844:	5'- CGCCTTTCCTTTTTCCTCCA
	SEQ ID No. 845:	5'- GCCCCGGTTTCCCGGCGTTA
5	SEQ ID No. 846:	5'- GCCGCCTTTCCTTTTTCCTC
	SEQ ID No. 847:	5'- TAGCCCCGGTTTCCCGGCGT
	SEQ ID No. 848:	5'- CCGGGTACCGTCAAGGCGCC
	SEQ ID No. 849:	5'- AAGCCGCCTTTCCTTTTTC
	SEQ ID No. 850:	5'- CCCC GGTTTCCCGGCGTTAT
10	SEQ ID No. 851:	5'- CCGGCGTTATCCCAGTCTTA
	SEQ ID No. 852:	5'- AGCCGCCTTTCCTTTTTCCT
	SEQ ID No. 853:	5'- CCGCCTTTCCTTTTTCCTCC
	SEQ ID No. 854:	5'- TTAGCCCCGGTTTCCCGGCG
	SEQ ID No. 855:	5'- CCCGGCGTTATCCCAGTCTT
15	SEQ ID No. 856:	5'- GCCGGGTACCGTCAAGGCGC
	SEQ ID No. 857:	5'- GGCCGGGTACCGTCAAGGCG
	SEQ ID No. 858:	5'- TCCCGGCGTTATCCCAGTCT
	SEQ ID No. 859:	5'- TGGCCGGGTACCGTCAAGGC
	SEQ ID No. 860:	5'- GAAGCCGCCTTTCCTTTTTC
20	SEQ ID No. 861:	5'- CCCGGTTTCCCGGCGTTATC
	SEQ ID No. 862:	5'- CGGCGTTATCCCAGTCTTAC
	SEQ ID No. 863:	5'- GGCGTTATCCCAGTCTTACA
	SEQ ID No. 864:	5'- GCGTTATCCCAGTCTTACAG
	SEQ ID No. 865:	5'- CGGGTACCGTCAAGGCGCCG
25	SEQ ID No. 866:	5'- ATTAGCCCCGGTTTCCCGGC
	SEQ ID No. 867:	5'- AAGGGGAAGGCCCTGTCTCC
	SEQ ID No. 868:	5'- GGCCCTGTCTCCAGGGAGGT
	SEQ ID No. 869:	5'- AGGCCCTGTCTCCAGGGAGG
	SEQ ID No. 870:	5'- AAGGCCCTGTCTCCAGGGAG
30	SEQ ID No. 871:	5'- GCCCTGTCTCCAGGGAGGTC

	SEQ ID No. 872:	5'- CGTTATCCCAGTCTTACAGG
	SEQ ID No. 873:	5'- GGGTACCGTCAAGGCGCCGC
	SEQ ID No. 874:	5'- CGGCAACAGAGTTTTACGAC
	SEQ ID No. 875:	5'- GGGGAAGGCCCTGTCTCCAG
5	SEQ ID No. 876:	5'- AGGGGAAGGCCCTGTCTCCA
	SEQ ID No. 877:	5'- GCAGCCGAAGCCGCCTTTCC
	SEQ ID No. 878:	5'- TTCTTCCCCGGCAACAGAGT
	SEQ ID No. 879:	5'- CGGCACTTGTTCTTCCCCGG
	SEQ ID No. 880:	5'- GTTCTTCCCCGGCAACAGAG
10	SEQ ID No. 881:	5'- GGCACCTGTTCTTCCCCGGC
	SEQ ID No. 882:	5'- GCACTTGTTCTTCCCCGGCA
	SEQ ID No. 883:	5'- CACTTGTTCTTCCCCGGCAA
	SEQ ID No. 884:	5'- TCTTCCCCGGCAACAGAGTT
	SEQ ID No. 885:	5'- TTGTTCTTCCCCGGCAACAG
15	SEQ ID No. 886:	5'- ACTTGTTCTTCCCCGGCAAC
	SEQ ID No. 887:	5'- TGTTCTTCCCCGGCAACAGA
	SEQ ID No. 888:	5'- CTTGTTCTTCCCCGGCAACA
	SEQ ID No. 889:	5'- ACGGCACTTGTTCTTCCCCG
	SEQ ID No. 890:	5'- GTCCGCCGCTAACCTTTTAA
20	SEQ ID No. 891:	5'- CTGGCCGGGTACCGTCAAGG
	SEQ ID No. 892:	5'- TCTGGCCGGGTACCGTCAAG
	SEQ ID No. 893:	5'- TTCTGGCCGGGTACCGTCAA
	SEQ ID No. 894:	5'- CAATGCTGGCAACTAAGGTC
	SEQ ID No. 895:	5'- CGTCCGCCGCTAACCTTTTA
25	SEQ ID No. 896:	5'- CGAAGCCGCCTTTCCTTTTT
	SEQ ID No. 897:	5'- CCGAAGCCGCCTTTCCTTTT
	SEQ ID No. 898:	5'- GCCGAAGCCGCCTTTCCTTT
	SEQ ID No. 899:	5'- AGCCGAAGCCGCCTTTCCTT
	SEQ ID No. 900:	5'- ACCGTCAAGGCGCCGCCCTG
30	SEQ ID No. 901:	5'- CCGTGGCTTTCTGGCCGGGT

SEQ ID No. 902: 5'- GCTTTCTGGCCGGGTACCGT  
SEQ ID No. 903: 5'- GCCGTGGCTTTCTGGCCGGG  
SEQ ID No. 904: 5'- GGCTTTCTGGCCGGGTACCG  
SEQ ID No. 905: 5'- CTTTCTGGCCGGGTACCGTC  
5 SEQ ID No. 906: 5'- TGGCTTTCTGGCCGGGTACC  
SEQ ID No. 907: 5'- GTGGCTTTCTGGCCGGGTAC  
SEQ ID No. 908: 5'- CGTGGCTTTCTGGCCGGGTAC  
SEQ ID No. 909: 5'- TTTCTGGCCGGGTACCGTCA  
SEQ ID No. 910: 5'- GGGAAGGCCCTGTCTCCAGG  
10 SEQ ID No. 911: 5'- CGAAGGGGAAGGCCCTGTCT  
SEQ ID No. 912: 5'- CCGAAGGGGAAGGCCCTGTC  
SEQ ID No. 913: 5'- GAAGGGGAAGGCCCTGTCTC  
SEQ ID No. 914: 5'- GGCGCCGCCCTGTTCGAACG  
SEQ ID No. 915: 5'- AGGCGCCGCCCTGTTCGAAC  
15 SEQ ID No. 916: 5'- AAGGCGCCGCCCTGTTCGAA  
SEQ ID No. 917: 5'- CCCGGCAACAGAGTTTTACG  
SEQ ID No. 918: 5'- CCCC GGCAACAGAGTTTTAC  
SEQ ID No. 919: 5'- CCATCTGTAAGTGGCAGCCG  
SEQ ID No. 920: 5'- TCTGTAAGTGGCAGCCGAAG  
20 SEQ ID No. 921: 5'- CTGTAAGTGGCAGCCGAAGC  
SEQ ID No. 922: 5'- CCCATCTGTAAGTGGCAGCC  
SEQ ID No. 923: 5'- TGTAAGTGGCAGCCGAAGCC  
SEQ ID No. 924: 5'- CATCTGTAAGTGGCAGCCGA  
SEQ ID No. 925: 5'- ATCTGTAAGTGGCAGCCGAA  
25 SEQ ID No. 926: 5'- CAGCCGAAGCCGCCTTTCCT  
SEQ ID No. 927: 5'- GGCAACAGAGTTTTACGACC  
SEQ ID No. 928: 5'- CCGGCAACAGAGTTTTACGA  
SEQ ID No. 929: 5'- TTCCCCGGCAACAGAGTTTT  
SEQ ID No. 930: 5'- CTTCCCCGGCAACAGAGTTT  
30 SEQ ID No. 931: 5'- TCCCCGGCAACAGAGTTTAA

	SEQ ID No. 932:	5'- CCGTCCGCCGCTAACCTTTT
	SEQ ID No. 933:	5'- CTTCTCCGACTTACGCCGG
	SEQ ID No. 934:	5'- CCTCCGACTTACGCCGGCAG
	SEQ ID No. 935:	5'- TTCCTCCGACTTACGCCGGC
5	SEQ ID No. 936:	5'- TCCTCCGACTTACGCCGGCA
	SEQ ID No. 937:	5'- TCCGACTTACGCCGGCAGTC
	SEQ ID No. 938:	5'- CCGACTTACGCCGGCAGTCA
	SEQ ID No. 939:	5'- GCCTTCCTCCGACTTACGCC
	SEQ ID No. 940:	5'- CCTTCCTCCGACTTACGCCG
10	SEQ ID No. 941:	5'- GCTCTCCCCGAGCAACAGAG
	SEQ ID No. 942:	5'- CTCTCCCCGAGCAACAGAGC
	SEQ ID No. 943:	5'- CGCTCTCCCCGAGCAACAGA
	SEQ ID No. 944:	5'- CTCCGACTTACGCCGGCAGT
	SEQ ID No. 945:	5'- TCTCCCCGAGCAACAGAGCT
15	SEQ ID No. 946:	5'- CGACTTACGCCGGCAGTCAC
	SEQ ID No. 947:	5'- TCGGCACTGGGGTGTGTCCC
	SEQ ID No. 948:	5'- GGCAC TGGGGTGTGTCCCCC
	SEQ ID No. 949:	5'- CTGGGGTGTGTCCCCCAAC
	SEQ ID No. 950:	5'- CACTGGGGTGTGTCCCCCA
20	SEQ ID No. 951:	5'- ACTGGGGTGTGTCCCCCAA
	SEQ ID No. 952:	5'- GCACTGGGGTGTGTCCCCC
	SEQ ID No. 953:	5'- TGGGGTGTGTCCCCCAACA
	SEQ ID No. 954:	5'- CACTCCAGACTTGCTCGACC
	SEQ ID No. 955:	5'- TCACTCCAGACTTGCTCGAC
25	SEQ ID No. 956:	5'- CGGCACTGGGGTGTGTCCCC
	SEQ ID No. 957:	5'- CGCCTTCCTCCGACTTACGC
	SEQ ID No. 958:	5'- CTCCCCGAGCAACAGAGCTT
	SEQ ID No. 959:	5'- ACTCCAGACTTGCTCGACCG
	SEQ ID No. 960:	5'- CCCATGCCGCTCTCCCCGAG
30	SEQ ID No. 961:	5'- CCATGCCGCTCTCCCCGAGC



SEQ ID No. 962: 5'- CCCCATGCCGCTCTCCCCGA  
SEQ ID No. 963: 5'- TCACTCGGTACCGTCTCGCA  
SEQ ID No. 964: 5'- CATGCCGCTCTCCCCGAGCA  
SEQ ID No. 965: 5'- ATGCCGCTCTCCCCGAGCAA  
5 SEQ ID No. 966: 5'- TTCGGCACTGGGGTGTGTCC  
SEQ ID No. 967: 5'- TGCCGCTCTCCCCGAGCAAC  
SEQ ID No. 968: 5'- TTCACTCCAGACTTGCTCGA  
SEQ ID No. 969: 5'- CCCGCAAGAAGATGCCTCCT  
SEQ ID No. 970: 5'- AGAAGATGCCTCCTCGCGGG  
10 SEQ ID No. 971: 5'- AAGAAGATGCCTCCTCGCGG  
SEQ ID No. 972: 5'- CGCAAGAAGATGCCTCCTCG  
SEQ ID No. 973: 5'- AAGATGCCTCCTCGCGGGCG  
SEQ ID No. 974: 5'- CCGCAAGAAGATGCCTCCTC  
SEQ ID No. 975: 5'- GAAGATGCCTCCTCGCGGGC  
15 SEQ ID No. 976: 5'- CCCC GCAAGAAGATGCCTCC  
SEQ ID No. 977: 5'- CAAGAAGATGCCTCCTCGCG  
SEQ ID No. 978: 5'- TCCTTCGGCACTGGGGTGTG  
SEQ ID No. 979: 5'- CCGCTCTCCCCGAGCAACAG  
SEQ ID No. 980: 5'- TGCCTCCTCGCGGGCGTATC  
20 SEQ ID No. 981: 5'- GACTTACGCCGGCAGTCACC  
SEQ ID No. 982: 5'- GGCTCCTCTCTCAGCGGGCC  
SEQ ID No. 983: 5'- CCTTCGGCACTGGGGTGTGT  
SEQ ID No. 984: 5'- GGGGTGTGTCCCCCAACAC  
SEQ ID No. 985: 5'- GCCGCTCTCCCCGAGCAACA  
25 SEQ ID No. 986: 5'- AGATGCCTCCTCGCGGGCGT  
SEQ ID No. 987: 5'- CACTCGGTACCGTCTCGCAT  
SEQ ID No. 988: 5'- CTC ACTCGGTACCGTCTCGC  
SEQ ID No. 989: 5'- GCAAGAAGATGCCTCCTCGC  
SEQ ID No. 990: 5'- CTCCAGACTTGCTCGACCGC  
30 SEQ ID No. 991: 5'- TTACGCCGGCAGTCACCTGT

SEQ ID No. 992: 5'- CTTCGGCACTGGGGTGTGTC  
SEQ ID No. 993: 5'- CTCGCGGGCGTATCCGGCAT  
SEQ ID No. 994: 5'- GCCTCCTCGCGGGCGTATCC  
SEQ ID No. 995: 5'- ACTCGGTACCGTCTCGCATG  
5 SEQ ID No. 996: 5'- GATGCCTCCTCGCGGGCGTA  
SEQ ID No. 997: 5'- GGGTGTGTCCCCCAACACC  
SEQ ID No. 998: 5'- ACTTACGCCGGCAGTCACCT  
SEQ ID No. 999: 5'- CTTACGCCGGCAGTCACCTG  
SEQ ID No. 1000: 5'- ATGCCTCCTCGCGGGCGTAT  
10 SEQ ID No. 1001: 5'- GCGCCGCGGGCTCCTCTCTC  
SEQ ID No. 1002: 5'- GGTGTGTCCCCCAACACCT  
SEQ ID No. 1003: 5'- GTGTGTCCCCCAACACCTA  
SEQ ID No. 1004: 5'- CCTCGCGGGCGTATCCGGCA  
SEQ ID No. 1005: 5'- CCTCACTCGGTACCGTCTCG  
15 SEQ ID No. 1006: 5'- TCCTCACTCGGTACCGTCTC  
SEQ ID No. 1007: 5'- TCGCGGGCGTATCCGGCATT  
SEQ ID No. 1008: 5'- TTCACTCCAGACTTGCTCG  
SEQ ID No. 1009: 5'- TACGCCGGCAGTCACCTGTG  
SEQ ID No. 1010: 5'- TCCAGACTTGCTCGACCGCC  
20 SEQ ID No. 1011: 5'- CTCGGTACCGTCTCGCATGG  
SEQ ID No. 1012: 5'- CGCGGGCGTATCCGGCATT  
SEQ ID No. 1013: 5'- GCGTATCCGGCATTAGCGCC  
SEQ ID No. 1014: 5'- GGGCTCCTCTCTCAGCGGCC  
SEQ ID No. 1015: 5'- TCCCCGAGCAACAGAGCTTT  
25 SEQ ID No. 1016: 5'- CCCCAGCAACAGAGCTTTA  
SEQ ID No. 1017: 5'- CCGAGCAACAGAGCTTTACA  
SEQ ID No. 1018: 5'- CCATCCCATGGTTGAGCCAT  
SEQ ID No. 1019: 5'- GTGTCCCCCAACACCTAGC  
SEQ ID No. 1020: 5'- GCGGGCGTATCCGGCATTAG  
30 SEQ ID No. 1021: 5'- CGAGCGGCTTTTTGGGTTTC

SEQ ID No. 1022: 5'- CTTTCACTCCAGACTTGCTC  
SEQ ID No. 1023: 5'- TTCCTTCGGCACTGGGGTGT  
SEQ ID No. 1024: 5'- CCGCCTTCCTCCGACTTACG  
SEQ ID No. 1025: 5'- CCCGCCTTCCTCCGACTTAC  
5 SEQ ID No. 1026: 5'- CCTCCTCGCGGGCGTATCCG  
SEQ ID No. 1027: 5'- TCCTCGCGGGCGTATCCGGC  
SEQ ID No. 1028: 5'- CATTAGCGCCCGTTTCCGGG  
SEQ ID No. 1029: 5'- GCATTAGCGCCCGTTTCCGG  
SEQ ID No. 1030: 5'- GGCATTAGCGCCCGTTTCCG  
10 SEQ ID No. 1031: 5'- GTCTCGCATGGGGCTTTCCA  
SEQ ID No. 1032: 5'- GCCATGGACTTTCACTCCAG  
SEQ ID No. 1033: 5'- CATGGACTTTCACTCCAGAC  
SEQ ID No. 1037: 5'- ACCGTCTCACAAGGAGCTTT  
SEQ ID No. 1038: 5'- TACCGTCTCACAAGGAGCTT  
15 SEQ ID No. 1039: 5'- GTACCGTCTCACAAGGAGCT  
SEQ ID No. 1040: 5'- GCCTACCCGTGTATTATCCG  
SEQ ID No. 1041: 5'- CCGTCTCACAAGGAGCTTTC  
SEQ ID No. 1042: 5'- CTACCCGTGTATTATCCGGC  
SEQ ID No. 1043: 5'- GGTACCGTCTCACAAGGAGC  
20 SEQ ID No. 1044: 5'- CGTCTCACAAGGAGCTTTCC  
SEQ ID No. 1045: 5'- TCTCACAAGGAGCTTTCCAC  
SEQ ID No. 1046: 5'- TACCCGTGTATTATCCGGCA  
SEQ ID No. 1047: 5'- GTCTCACAAGGAGCTTTCCA  
SEQ ID No. 1048: 5'- ACCCGTGTATTATCCGGCAT  
25 SEQ ID No. 1049: 5'- CTCGGTACCGTCTCACAAGG  
SEQ ID No. 1050: 5'- CGGTACCGTCTCACAAGGAG  
SEQ ID No. 1051: 5'- ACTCGGTACCGTCTCACAAG  
SEQ ID No. 1052: 5'- CGGCTGGCTCCATAACGGTT  
SEQ ID No. 1053: 5'- ACAAGTAGATGCCTACCCGT  
30 SEQ ID No. 1054: 5'- TGGCTCCATAACGGTTACCT

SEQ ID No. 1055: 5'- CAAGTAGATGCCTACCCGTG  
SEQ ID No. 1056: 5'- CACAAGTAGATGCCTACCCG  
SEQ ID No. 1057: 5'- GGCTCCATAACGGTTACCTC  
SEQ ID No. 1058: 5'- ACACAAGTAGATGCCTACCC  
5 SEQ ID No. 1059: 5'- CTGGCTCCATAACGGTTACC  
SEQ ID No. 1060: 5'- GCTGGCTCCATAACGGTTAC  
SEQ ID No. 1061: 5'- GGCTGGCTCCATAACGGTTA  
SEQ ID No. 1062: 5'- GCTCCATAACGGTTACCTCA  
SEQ ID No. 1063: 5'- AAGTAGATGCCTACCCGTGT  
10 SEQ ID No. 1064: 5'- CTCCATAACGGTTACCTCAC  
SEQ ID No. 1065: 5'- TGCCTACCCGTGTATTATCC  
SEQ ID No. 1066: 5'- TCGGTACCGTCTCACAAGGA  
SEQ ID No. 1067: 5'- CTCACAAGGAGCTTTCCACT  
SEQ ID No. 1068: 5'- GTAGATGCCTACCCGTGTAT  
15 SEQ ID No. 1069: 5'- CCTACCCGTGTATTATCCGG  
SEQ ID No. 1070: 5'- CACTCGGTACCGTCTCACAA  
SEQ ID No. 1071: 5'- CTCAGCGATGCAGTTGCATC  
SEQ ID No. 1072: 5'- AGTAGATGCCTACCCGTGTA  
SEQ ID No. 1073: 5'- GCGGCTGGCTCCATAACGGT  
20 SEQ ID No. 1074: 5'- CCAAAGCAATCCCAAGGTTG  
SEQ ID No. 1075: 5'- TCCATAACGGTTACCTCACC  
SEQ ID No. 1076: 5'- CCCGTGTATTATCCGGCATT  
SEQ ID No. 1077: 5'- TCTCAGCGATGCAGTTGCAT  
SEQ ID No. 1078: 5'- CCATAACGGTTACCTCACCG  
25 SEQ ID No. 1079: 5'- TCAGCGATGCAGTTGCATCT  
SEQ ID No. 1080: 5'- GGCGGCTGGCTCCATAACGG  
SEQ ID No. 1081: 5'- AAGCAATCCCAAGGTTGAGC  
SEQ ID No. 1082: 5'- TCACTCGGTACCGTCTCACA  
SEQ ID No. 1083: 5'- CCGAGTGTTATTCCAGTCTG  
30 SEQ ID No. 1084: 5'- CACAAGGAGCTTTCCACTCT

SEQ ID No. 1085: 5'- ACAAGGAGCTTTCCACTCTC  
SEQ ID No. 1086: 5'- TCACAAGGAGCTTTCCACTC  
SEQ ID No. 1087: 5'- CAGCGATGCAGTTGCATCTT  
SEQ ID No. 1088: 5'- CAAGGAGCTTTCCACTCTCC  
5 SEQ ID No. 1089: 5'- CCAGTCTGAAAGGCAGATTG  
SEQ ID No. 1090: 5'- CAGTCTGAAAGGCAGATTGC  
SEQ ID No. 1091: 5'- CGGCGGCTGGCTCCATAACG  
SEQ ID No. 1092: 5'- CCTCTCTCAGCGATGCAGTT  
SEQ ID No. 1093: 5'- CTCTCTCAGCGATGCAGTTG  
10 SEQ ID No. 1094: 5'- TCTCTCAGCGATGCAGTTGC  
SEQ ID No. 1095: 5'- CTCTCAGCGATGCAGTTGCA  
SEQ ID No. 1096: 5'- CAATCCCAAGGTTGAGCCTT  
SEQ ID No. 1097: 5'- AATCCCAAGGTTGAGCCTTG  
SEQ ID No. 1098: 5'- AGCAATCCCAAGGTTGAGCC  
15 SEQ ID No. 1099: 5'- CTCACTCGGTACCGTCTCAC  
SEQ ID No. 1100: 5'- GCAATCCCAAGGTTGAGCCT  
SEQ ID No. 1101: 5'- GCCTTGGA CTTTCACTTCAG  
SEQ ID No. 1102: 5'- CATAACGGTTACCTCACCGA  
SEQ ID No. 1103: 5'- CTCCTCTCTCAGCGATGCAG  
20 SEQ ID No. 1104: 5'- TCGGCGGCTGGCTCCATAAC  
SEQ ID No. 1105: 5'- AGTCTGAAAGGCAGATTGCC  
SEQ ID No. 1106: 5'- TCCTCTCTCAGCGATGCAGT  
SEQ ID No. 1107: 5'- CCCAAGGTTGAGCCTTGGAC  
SEQ ID No. 1108: 5'- ATAACGGTTACCTCACCGAC  
25 SEQ ID No. 1109: 5'- TCCCAAGGTTGAGCCTTGGA  
SEQ ID No. 1110: 5'- ATTATCCGGCATTAGCACCC  
SEQ ID No. 1111: 5'- CTACGTGCTGGTAACACAGA  
SEQ ID No. 1112: 5'- GCCGCTAGCCCCGAAGGGCT  
SEQ ID No. 1113: 5'- CTAGCCCCGAAGGGCTCGCT  
30 SEQ ID No. 1114: 5'- CGCTAGCCCCGAAGGGCTCG

SEQ ID No. 1115: 5'- AGCCCCGAAGGGCTCGCTCG  
SEQ ID No. 1116: 5'- CCGCTAGCCCCGAAGGGCTC  
SEQ ID No. 1117: 5'- TAGCCCCGAAGGGCTCGCTC  
SEQ ID No. 1118: 5'- GCTAGCCCCGAAGGGCTCGC  
5 SEQ ID No. 1119: 5'- GCCCCGAAGGGCTCGCTCGA  
SEQ ID No. 1120: 5'- ATCCCAAGGTTGAGCCTTGG  
SEQ ID No. 1121: 5'- GAGCCTTGGACTTTCACTTC  
SEQ ID No. 1122: 5'- CAAGGTTGAGCCTTGGACTT  
SEQ ID No. 1123: 5'- GAGCTTTCCACTCTCCTTGT  
10 SEQ ID No. 1124: 5'- CCAAGGTTGAGCCTTGGACT  
SEQ ID No. 1125: 5'- CGGGCTCCTCTCTCAGCGAT  
SEQ ID No. 1126: 5'- GGAGCTTTCCACTCTCCTTG  
SEQ ID No. 1127: 5'- GGGCTCCTCTCTCAGCGATG  
SEQ ID No. 1128: 5'- TCTCCTTGTCGCTCTCCCCG  
15 SEQ ID No. 1129: 5'- TCCTTGTCGCTCTCCCCGAG  
SEQ ID No. 1130: 5'- AGCTTTCCACTCTCCTTGTC  
SEQ ID No. 1131: 5'- CCACTCTCCTTGTCGCTCTC  
SEQ ID No. 1132: 5'- GGCTCCTCTCTCAGCGATGC  
SEQ ID No. 1133: 5'- CCTTGTCGCTCTCCCCGAGC  
20 SEQ ID No. 1134: 5'- CACTCTCCTTGTCGCTCTCC  
SEQ ID No. 1135: 5'- ACTCTCCTTGTCGCTCTCCC  
SEQ ID No. 1136: 5'- CTCTCCTTGTCGCTCTCCCC  
SEQ ID No. 1137: 5'- GCGGGCTCCTCTCTCAGCGA  
SEQ ID No. 1138: 5'- GGCTCCATCATGGTTACCTC  
25 SEQ ID No. 1142: 5'- CTTCTCCGGCTTGCGCCGG  
SEQ ID No. 1143: 5'- CGCTCTTCCCGA(G/T)TGACTGA  
SEQ ID No. 1144: 5'- CCTCGGGCTCCTCCATC(A/T)GC

2. The method according to claim 1, wherein drink-spoiling microorganisms belonging to the genus *Zygosacchaeromyces* are detected with oligonucleotide probe SEQ ID No. 1.

5           3. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces bailii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 5 to SEQ ID No. 21.

10           4. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces fermentati* is detected with oligonucleotide probe SEQ ID No. 22.

15           5. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces microellipsoides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 23 to SEQ ID No. 24.

20           6. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces mellis* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 25 to SEQ ID No. 75.

25           7. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces rouxii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 76 to SEQ ID No. 126.

8. The method according to claim 1, wherein the drink-spoiling microorganisms *Zygosacchaeromyces mellis* and *Zygosacchaeromyces rouxii* are detected simultaneously with oligonucleotide probe SEQ ID No. 127.

5           9. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces bisporus* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 128 to SEQ ID No. 142.

10           10. The method according to claim 1, wherein the drink-spoiling microorganism *Hanseniaspora uvarum* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 143 to SEQ ID No. 144.

15           11. The method according to claim 1, wherein the drink-spoiling microorganism *Candida intermedia* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 145 to SEQ ID No. 146.

20           12. The method according to claim 1, wherein the drink-spoiling microorganism *Candida parapsilosis* is detected with oligonucleotide probe SEQ ID No. 148.

25           13. The method according to claim 1, wherein the drink-spoiling microorganism *Candida crusei* (*Issatchenkia orientalis*) is detected with oligonucleotide probe SEQ ID No. 149.

          14. The method according to claim 1, wherein the drink-spoiling microorganisms *Brettanomyces* (*Dekkera*) *anomala* and *Dekkera bruxellensis* are detected simultaneously with oligonucleotide probe SEQ ID No. 150.



15. The method according to claim 1, wherein the drink-spoiling microorganism *Brettanomyces (Dekkera) bruxellensis* is detected with oligonucleotide probe SEQ ID No. 151.

5           16. The method according to claim 1, wherein the drink-spoiling microorganism *Brettanomyces (Dekkera) naardenensis* is detected with oligonucleotide probe SEQ ID No. 152.

10           17. The method according to claim 1, wherein the drink-spoiling microorganism *Pichia membranaefaciens* is detected with oligonucleotide probe SEQ ID No. 153.

15           18. The method according to claim 1, wherein the drink-spoiling microorganisms *Pichia minuta* and *Pichia anomala* are detected simultaneously with oligonucleotide probe SEQ ID No. 154.

20           19. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomyces exiguus* is detected with oligonucleotide probe SEQ ID No. 157.

25           20. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomycodes ludwigii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 158 to SEQ ID No. 159.

21. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomyces cerevisiae* is detected with oligonucleotide probe SEQ ID No. 160.

22. The method according to claim 1, wherein the drink-spoiling microorganism *Mucor racemosus* is detected with oligonucleotide probe SEQ ID No. 163.

5           23. The method according to claim 1, wherein the drink-spoiling microorganism *Byssoschlamys nivea* is detected with oligonucleotide probe SEQ ID No. 164.

24. The method according to claim 1, wherein the drink-spoiling  
10 microorganism *Neosartorya fischeri* is detected with oligonucleotide probe SEQ ID No. 165.

25. The method according to claim 1, wherein the drink-spoiling microorganisms *Aspergillus fumigatus* and *A. fischeri* are detected simultaneously  
15 with oligonucleotide probe SEQ ID No. 166.

26. The method according to claim 1, wherein the drink-spoiling microorganism *Talaromyces flavus* is detected with oligonucleotide probe SEQ ID  
20 No. 167.

27. The method according to claim 1, wherein the drink-spoiling microorganisms *Talaromyces bacillisporus* and *T. flavus* are detected simultaneously  
with oligonucleotide probe SEQ ID No. 168.

25           28. The method according to claim 1, wherein the drink-spoiling microorganism *Lactobacillus collinoides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 169 to SEQ ID No. 269.

29. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Leuconostoc* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 270 to SEQ ID No. 271.

5           30. The method according to claim 1, wherein the drink-spoiling microorganisms *Leuconostoc mesenteroides* and *L. pseudomesenteroides* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 272 to SEQ ID No. 301.

10           31. The method according to claim 1, wherein the drink-spoiling microorganism *Leuconostoc pseudomesenteroides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 302 to SEQ ID No. 341.

15           32. The method according to claim 1, wherein the drink-spoiling microorganism *Oenococcus oenis* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 342 to SEQ ID No. 444.

20           33. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Weissella* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 445 to SEQ ID No. 495.

25           34. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Lactococcus* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 496 to SEQ ID No. 546.

30           35. The method according to claim 1, wherein drink-spoiling microorganisms of the genera *Acetobacter* and *Gluconobacter* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 547 to SEQ ID No. 608.

36. The method according to claim 1, wherein drink-spoiling microorganisms of the genera *Acetobacter*, *Gluconobacter* and *Gluconoacetobacter* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 609 to SEQ ID No. 842.

37. The method according to claim 1, wherein the drink-spoiling microorganism *Bacillus coagulans* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 843 to SEQ ID No. 932.

38. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Alicyclobacillus* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 933 to SEQ ID No. 1033.

39. The method according to claim 1, wherein the drink-spoiling microorganism *Alicyclobacillus acidoterrestris* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 1037 to SEQ ID No. 1138.

40. The method according to claim 1, wherein the drink-spoiling microorganisms *Alicyclobacillus cycloheptanicus* and *A. herbarius* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 1142 to SEQ ID No. 1144.

41. The method according to claim 2, characterised in that the at least one oligonucleotide probe is used in combination with one or more competitor probes.

42. The method according to claim 41, characterised in that the oligonucleotide probe SEQ ID No. 1 is used in combination with one or more

competitor probes selected from the group consisting of SEQ ID No. 2 to SEQ ID No. 4.

43. The method according to claim 11, characterised in that the at least one  
5 oligonucleotide probe is used in combination with one or more competitor probes.

44. The method according to claim 43, characterised in that the  
oligonucleotide probe SEQ ID No. 146 is used in combination with competitor probe  
SEQ ID No. 147.

10

45. The method according to claim 18, characterised in that the at least one  
oligonucleotide probe is used in combination with one or more competitor probes.

46. The method according to claim 45, characterised in that the  
15 oligonucleotide probe SEQ ID No. 154 is used in combination with one or more  
competitor probes selected from the group consisting of SEQ ID No. 155 to SEQ ID  
No. 156.

47. The method according to claim 21, characterised in that the at least one  
20 oligonucleotide probe is used in combination with one or more competitor probes.

48. The method according to claim 47, characterised in that the  
oligonucleotide probe SEQ ID No. 160 is used in combination with one or more  
competitor probes selected from the group consisting of SEQ ID No. 161 to SEQ ID  
25 No. 162.

49. The method according to claim 38, characterised in that the at least one  
oligonucleotide probe is used in combination with one or more competitor probes.

50. The method according to claim 49, characterised in that the oligonucleotide probe SEQ ID No. 933 is used in combination with one or more competitor probes selected from the group consisting of SEQ ID No. 1034 to SEQ ID No. 1036.

5

51. The method according to claim 39, characterised in that the at least one oligonucleotide probe is used in combination with one or more competitor probes.

52. The method according to claim 51, characterised in that the oligonucleotide probe SEQ ID No. 1044 is used in combination with the competitor probe SEQ ID No. 1139.

53. The method according to claim 51, characterised in that the oligonucleotide probe SEQ ID No. 1057 is used in combination with one or more competitor probes selected from the group consisting of SEQ ID No. 1140 to SEQ ID No. 1141.

54. The method according to any of claims 1 to 53, characterized in by comprising the following steps:

- 20
- a) cultivating the drink-spoiling microorganisms contained in the sample,
  - b) fixing the drink-spoiling microorganisms contained in the sample,
  - c) incubating the fixed microorganisms with at least one oligonucleotide probe optionally in combination with a competitor probe,
  - d) removing non-hybridised oligonucleotide probes,
  - 25 e) detecting and visualizing and optionally quantifying the drink-spoiling microorganisms with the hybridized oligonucleotide probes.

55. The method according to any of claims 1 to 54, characterized in that the sample is a sample from non-alcoholic beverages.

30

56. A kit for performing a method according to any of claims 1 to 55, containing at least one oligonucleotide according to claim 1.